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The National Academy of Medical Sciences (India) completed its 50 glorious years in 2011. Established as a unique institution which fosters and utilizes academic excellence as its resource to meet medical and social goals, the Academy has been recognised by the Government of India as a Nodal Agency for Continuing Education for Medical and Allied Health Professionals and is advising the Government of India in matters of National Health Policy and Planning.

The Academy encourages and sponsors nation-wide CME programmes, Symposia, Workshops and Conferences. Over the years the Academy has recognized the outstanding achievements made by the Indian scientists in the field of medicine and allied sciences and conferred Fellowship as well as Membership to the selected persons through a structured peer reviewed process and voting by all the Fellows.

One of the key role of Academy is dissemination of peer reviewed scientific material contributed by biomedical scientists. Efforts of our esteemed fellows have kept the literary and scientific contents of the Journal “Annals of the National Academy of Medical Sciences (India)” to the highest level. With unrelenting efforts of our dynamic community we have been able to maintain the continuity of the publication also. The Academy is trying to get the Annals indexed in PubMed and are leaving no stone unturned to get the job done. The major criteria for the same are the quality of articles being submitted and published, peer-review process, and regularity of publication. In its endeavour to do so, we are changing the size of the journal at par with our global counterparts. The Annals Vol 53, first issue is in your hands with improved size. Secondly, our Annals is now online also at annals-nams.in and all articles are available with open access. Not only the site provides access to reading, one can also submit their research and other contribution online without any charges. Thirdly, we have already started our indexing with various agencies to get wider coverage of our contents. The Annals is indexed with Index Medicus of South-East Asia region (IMSEAR) of WHO and is extracted by Directory of Open Access Journal (DOAJ), Google Scholar, Harvard Database, BASE, Citefactor and other indexing agencies.

The family of our Academy has grown enormously in size in numbers, variety and multidisciplinary talent. We have more than 900 Fellows and 6300 members as of December 2016. Whenever we wish to share our scientific achievements, we tend to become very selective in choosing the journal and our criteria are restricted on indexing status, suitability for the journal and type of research work focused on particular speciality and journal impact factor. With the support given by all fellows and members, Annals is now getting manuscripts which are aligned with our mandate to uplift the status of public health, medical education with emphasis on competency and skill development, and innovations in healthcare. One of the unique quality of Annals is its diversity and assortment rarely seen in speciality journals. The current issue exemplify this diversity. Readers gets an opportunity to peep into domain of other scientists who are doing path breaking work in their areas of expertise.

Many epigenetic and genetic factors may determine diabetic patients’ susceptibility to renal disease
development. Diabetic Nephropathy (DN) is a result of multifactorial mechanisms and may ultimately lead to Chronic Kidney Disease (CKD). Dr O P Kalra has shared his work on "Genetic Basis of Diabetic Nephropathy" based on his extensive experience in this issue of the Annals.

For a long time Medical Educationists in India are struggling to implement reforms in curriculum both for under-graduates and post-graduates education. Many radical changes have been implemented in curriculum in Western world and but are still being evaluated for their learning outcomes. No better time exists for medical faculty in India to just carry out smaller Quality Improvement (QI) changes in existing system and study the impact in existing settings where Indian system of education is still producing physician who are being recognized and appreciated for their skills globally as the world is closely watching the developments in medical education in India. Post-graduate students’ self-assessment regarding their competencies and skills showed adequacy as far as Communication and health education is concerned but lacked confidence in epidemiology and occupational health. This was the conclusion drawn in a study by Kishore et al. Authors felt a need for reforms in existing curriculum.

Innovations do help mankind in preserving their health has been immensely proved by work by Dr. Gulati and Dr. Dash. Despite improvement in technology and advances in healthcare, we are still not able to achieve a single digit score in Infant Mortality Rate except in state of Kerala. To a large extent perinatal factors and genetic conditions are responsible among survivors having adverse developmental outcome and disability. The children need an early detection of neurodevelopmental conditions and early intervention coupled with augmentation of skills among the healthcare workers at grass root level. This can be achieved by developing pediatric sub-specialities with innovative approaches towards diagnosis, management and research. Autism was once a rare diagnosis. Observant paediatricians are now picking more cases from their busy practices coupled with the help of informed public. Innovative tools have been widely researched in India and are making life of the families having child suffering from Autistic Spectrum Disorder (ASD) much easier. All credit goes to Indian biomedical scientists who selflessly get involved with these children and their families. Dr. Gulati from AIIMS Delhi has shared her dream of developing Pediatric Neurology and how the vision got realized through offbeat journey less travelled by others is being highlighted in her article “Neurodevelopmental disorders: The Journey, the dreams and their realization”. On the other hand Dr. Dash from BHU, Varanasi has described innovative applications of Nanomaterials. Graphene based biosensor can detect individuals with high risk for thrombosis while near-infrared laser-irradiated gold Nano rods can ablate pathologic thrombus in-situ. Academy provides a platform for biomedical scientists for sharing, networking and integrating all medical specialities since the collaboration is the buzzword for all round development of health of mankind.

Public awareness with the help of mass media, mobile technology and education has definitely brought down diseases related to infections, poverty and nutritional disorders to great extent if not completely. However, India is facing problems with non-communicable diseases and emerging infections due to changes in socio-environmental milieu and changing lifestyles. The lurking danger of cancer, metabolic disorders and silent diseases like hypertension and diabetes mellitus on one hand and continuing threats of infectious diseases on the other hand is creating heavy stress on our healthcare system. Unless we take our vision to a futuristic horizon and start harnessing technology coupled with clinical research we will be at great disadvantage in the healthcare we are going to provide. Neuro-
restoration in stroke patients using multimodality approaches incorporating stem cells, robotics and drugs do help patients regain their functional capabilities has been highlighted by Dr. Padma Srivastava. She has delved deeper into newer cell based therapies, appliances, drugs and devices which have worked advantageously both in experimental and clinical studies in stroke patients.

Editorial board hopes that this present issue will instil new life in our journey towards academic excellence in showcasing the admirable work of Indian biomedical scientists to our global community.

Dr. Sanjeev Misra
Genetic Basis of Diabetic Nephropathy

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ABSTRACT

It is well known that all patients with Type 2 Diabetes Mellitus (T2DM) do not develop chronic kidney disease (CKD). Several metabolic, hemodynamic and intracellular mechanisms have been proposed to play role in the pathogenesis of Diabetic Nephropathy (DN). Clustering of patients with DN in certain ethnic groups and families suggests the role of genetic factors. We have studied various facets about genetic determinants which may influence the development of kidney disease in patients with T2DM.

We have found that Angiotensin Converting Enzyme (ACE) DD genotype conferred the maximum risk, whereas ACE II genotype seemed to confer protective role against development of diabetic and nondiabetic CKD. Further, we found that oxidative stress (OS) plays a significant role in the development of DN and that Glutathione S-transferase theta-1 and/or Glutathione S-transferase Mu-1 null genotypes are associated with higher OS in patients with DN. In addition, we also found that increased levels of inflammatory mediators, i.e. Tumor necrosis factor-α (TNF-α), high-sensitivity C-reactive protein (hsCRP) and Urinary Monocyte Chemoattractant Protein-1 (uMCP-1) play a significant role in contributing to OS. We have shown that genetic polymorphism of NF-kB gene and TNF-α gene plays a role in determining serum level of various inflammatory markers and oxidant stress parameters. We found significant association of -429T/C and Gly82Ser Receptors for Advanced Glycation End-products (RAGE) polymorphisms with the development of macrovascular and microvascular complications, respectively in T2DM subjects. Further, we have observed that AGE-mediated exacerbation of RAGE expression may play a role in pathogenesis of various vascular complications in T2DM.

To conclude, polymorphisms of various genes involved in renin-angiotensin aldosterone system (RAAS), inflammatory, oxidant stress, cytoprotective and nitrous oxide pathways and enhanced RAGE mRNA expression may adversely influence final common pathway through oxidant stress mechanisms, and influence the levels of various cytokines and intracellular signaling mechanisms, thereby influencing the susceptibility of patients with diabetes mellitus for development of kidney disease and vascular complications.

Keywords: Diabetic nephropathy, genetic factor, chronic disease.
Introduction

Diabetes Mellitus (DM) has become a leading cause of morbidity and mortality and is considered a major public health problem that places a significant burden on global healthcare resources. It is, however, much less appreciated that the diabetes epidemic would also be accompanied by an epidemic of Chronic Kidney Disease (CKD), which brings with it a huge burden of Cardiovascular Disease (CVD) and End-stage Renal Disease (ESRD), leading to premature death. It is estimated that the number of people with diabetes will rise from 171 million in 2000 to 366 million in 2030, resulting in millions of new cases of CKD, most of them being in the developing world (1). Approximately one-fourth to one-third of all diabetics go on to develop Diabetic Nephropathy (DN), making it one of the leading cause of CKD and ESRD requiring renal replacement therapy (2,3). Recent data from CKD Registry of India shows that DN accounts for the single largest group of patients with CKD (4). Since treating ESRD is simply unaffordable for most developing countries, the emphasis has to be on prevention, early detection and slowing of progression from early stages of CKD to ESRD.

Despite relentless research, the complex etiopathogenesis of kidney disease in DM has not been fully understood. The pathogenesis of DN is clearly multifactorial and several metabolic, hemodynamic and intracellular mechanisms or factors have been proposed to play a role in the onset and progression of disease (5), and are currently under active investigation at various centres all over the world. In addition, certain hereditary and environmental factors have been implicated in the etiopathogenesis of DN. During the last few years, we have focused our research into various genetic factors which may be potentially associated with the development of nephropathy in patients with Type 2 diabetes mellitus (T2DM).

Predictors for Development of Nephropathy in DM

There are various factors which play a role in the development of nephropathy in DM. The most well-known amongst these include poor glycemic control, family history of diabetes or hypertension, increased activity of sodium-lithium counter-transport mechanism in red blood cells, etc. Of these, most of the studies have focused on the role of glycemic control in the development of various complications. Studies done in Type 1 diabetes mellitus (T1DM), such as Diabetes Control and Complications Trial (DCCT) have shown that tight glycemic control using multiple insulin injections reduces the incidence of microalbuminuria by 39% (6). Similarly, in United Kingdom Prospective Diabetes Study (UKPDS), a 30% risk reduction for the development of microalbuminuria was observed in the group intensively treated for hyperglycemia (7). In the Kumamoto Study, intensive glycemic control reduced the rate of micro- and macroalbuminuria (8). However, it is pertinent to note that in these studies several patients developed DN despite tight glycemic control and vice versa. In a cross-sectional survey of 507 patients with T2DM, we found that self-efficacy was the single most important determinant of current diabetes control (p<0.01) and self-efficacy was influenced by various factors, such as educational status, employment, family support and mental attitude (9). Previously, it was believed that once albuminuria had become persistent, glycemic control lost its beneficial effect on the kidney, but several recent studies have documented the importance of glycemic control on the progression of nephropathy in patients with T1DM. Among the most important putative promoters of progression of kidney disease, blood pressure has been documented to have a close relation with rate of decline of glomerular filtration rate in both T1DM and T2DM. Serum cholesterol concentration has been shown to be another predictor of progression of nephropathy in both types of diabetes. Dietary protein
restriction retards the progression of renal disease in diabetes while smoking has been suggested to play a role in the progression of nephropathy in both types of diabetes.

**Hereditary and Ethnic Factors**

Present day knowledge states that specific genetic backgrounds might influence the development of DN. Indeed, only 30% patients with T1DM and 40% patients with T2DM develop DN irrespective of treatment for diabetes (10), and DN often shows a familial clustering in siblings with diabetes. It has been noted that the prevalence rates of DN in subjects with T2DM show a marked ethnic variation. Higher rates of diabetic renal disease are seen in Indo-Asians in the UK, in African-Americans (11), in Nauruans (12) and Pima Indians (13). The reason for this inter-racial difference in the incidence of DN is unclear, but ethnic variation in genetic susceptibility to nephropathy is a possibility. It is noteworthy that these ethnic groups not only have a very high incidence of T2DM, but also a high incidence of hypertension. This suggests that differences in genetic predisposition to hypertension may contribute to the higher prevalence of nephropathy in certain racial groups; although an alternative explanation may be that the presence of hypertension may accelerate an already present renal disease and lead to the condition becoming clinically manifest more quickly.

**Rationale for Genetic Studies in DN**

The fact that, a fairly large number of patients with DM goes on to develop nephropathy even in the absence of various factors mentioned above, has led scientists to postulate and investigate various genetic factors leading to this dreadful complication. There is enough evidence supporting the concept of genetic susceptibility to nephropathy in patients with diabetes (14,15). Discovery of genetic variants that underpin susceptibility to nephropathy could yield important insights into this condition. Firstly, it would permit identification of patients at risk of nephropathy shortly after diagnosis of diabetes rather than much later when persistent microalbuminuria develops, by which time there is already histological evidence of renal injury. This would facilitate targeted therapeutic interventions aimed at primary prevention rather than secondary treatment of established nephropathy. Secondly, and perhaps more importantly, if the susceptibility variants are located in genes that have not previously been implicated in DN, this may lead to improved understanding of its pathophysiology and development of targeted novel therapies.

**Strategies for Identifying Susceptibility Genes**

The etiology of DN is multifactorial, yet there is clear evidence of genetic basis. Strong association of familial aggregation and the heritability of DN in patients with T2DM provide compelling evidence that DN and its related traits are influenced by genetic factors and suggest a complex, multifactorial mode of inheritance with one or more major susceptibility genes. Familial clustering of renal disease in T2DM supports the hypothesis that the increased risk of DN in T2DM is partly due to a shared gene or set of genes among affected family members and has motivated investigations aimed at identifying the specific chromosomal regions that harbor genes contributing to its susceptibility. The major approaches that are currently being used to identify DN susceptibility genes are: (i) Candidate gene approach, (ii) Linkage analysis, and (iii) Genome-wide association studies.

**The Candidate Gene Approach**

The candidate gene approach involves assessment of genetic variation, typically Single Nucleotide Polymorphisms (SNPs) in one or more genes with plausible physiological roles in DN. These SNPs lie within a candidate gene or
region and are selected from the literature or from the Hap Map database (www.hapmap.org). The goal is to demonstrate a significant difference in allele frequencies between cases with DN and control subjects. In various studies done in our laboratory at University College of Medical Sciences, Delhi, in the field of DN, we have followed the candidate gene approach. A large number of candidate genes involved in several pathways have been tested for association with the development and progression of DN based on the possible physiological role of the genes in patients with DM and kidney disease, such as, renin-angiotensin aldosterone system (RAAS), glucose metabolism, growth factors, oxidative stress (OS), inflammation, lipid metabolism, etc.

Several candidate gene studies involving the above-mentioned genes to study the association with DN have been reported; however, the results have largely been inconsistent. Limitations of this approach include that candidate gene studies are frequently based on small number of cases and controls resulting in underpowered analyses. Various meta-analyses are being carried out to overcome the limitations of individual candidate gene study. Various candidate genes which we have studied belonging to different classes are discussed below:

A. Genes Involved in RAAS

RAAS has been shown to play a central role in the pathogenesis of most forms of CKD. Prorenin, renin, Angiotensin-Converting Enzyme (ACE) and angiotensin II levels are all noted to be elevated in DN (16). Furthermore, genes of the RAAS have been suggested as being genetic determinants for both hypertension and CVD, both of which are common in patients with DN.

Polymorphism of ACE gene has been implicated in determining the blood level of ACE and thereby may play an important role in the pathogenesis of DN. Initially we did a pilot study to investigate the prevalence of polymorphisms of ACE genotype in 100 subjects including patients with T2DM with/without DN, patients with non-diabetic CKD and healthy controls (17). We found that D allele of ACE gene acts as a risk factor for the development of nephropathy in patients with T2DM as well as for nondiabetic CKD, while I allele of ACE gene was protective in nature.

Few other investigators have studied ACE gene insertion-deletion polymorphism in DN patients with T2DM (18). A large meta-analysis found the association of ACE D allele with DN risk with an odds ratio (OR) in the range of 1.25–1.57 in the Asian subgroup (19); however, no significant effects were detected for the Caucasian subgroup. In DN, two small studies have suggested an association between the D allele of the ACE gene and nephropathy (20,21); however, other subsequent large studies with and without nephropathy have shown no association between nephropathy and the D allele (22). Overall, the cumulative results from a large number of studies suggest that if the ACE gene has any effect, it is likely to be small, and it is not useful as a screening marker for nephropathy.

In an ongoing study on 'Role of genetic polymorphisms of RAAS on the reno-protective efficacy of ACE inhibitors in patients with DN', we are studying various genes involved in RAAS, viz., ACE (I/D), angiotensinogen (AGT M235T) and angiotensin type I receptor genes in 255 patients of T2DM with nephropathy. Genotype frequency for ACE I/D polymorphism was II-36.7%, ID-50.0%, DD-13.3% and allele frequency was found to be I-61.7% and D-38.3%. We found that ACE inhibitor treatment in patients with DN resulted in significant reduction in urinary protein excretion which was found to be independent of ACE I/D and AGT M235T polymorphism (23).

All the patients have been put on ACE inhibitor therapy and are being followed-up at three-month intervals for a period of minimum
two years along with monitoring of albumin: creatinine ratio and estimated GFR to assess whether the gene polymorphism of RAAS can modulate the degree of beneficial response seen following ACE inhibitor therapy in preventing the progression of DN.

B. Genes Involved in Inflammatory Pathways

Association of biomarkers of inflammation with the risk of chronic kidney disease in T2DM

Traditionally, DN has been considered a nonimmune, degenerative disease; however, in 1991, Bohle et al (24) described the presence of monocytes, macrophages, T-cells, and fibroblasts associated with the tubulo-interstitial changes seen in DN. More recent reports (25, 26) have suggested that inflammation may underlie disease progression in DN. The activation of Nuclear Factor kappa B (NF-kB) - linked regulatory pathway generally underlies inflammatory processes, and an increase in the nuclear translocation of NF-kB has been demonstrated in human DN (27, 28). Polymorphism of NF-kB1 gene may influence activation / inactivation of NF-kB1 in renal cells which may influence urinary monocyte chemoattractant protein-1 (uMCP-1) levels in patients with DM (29, 30).

Recent evidence has highlighted the role of uMCP-1 in DN and showed it as a major factor influencing macrophage accumulation in renal disease. MCP-1 is a member of the CC chemokine family which is produced by endothelial cells, vascular smooth cells, keratinocytes, fibroblasts, mesangial cells, tubular epithelial cells, lymphocytes and monocytes/macrophages in response to a variety of pro-inflammatory stimuli. It is the strongest known chemotactic factor for monocytes and is upregulated in DN. Its expression has been identified in kidney diseases which involve significant inflammation (31-34). We recruited 150 subjects which were divided into 3 groups having 50 subjects in each group, viz; Group: I- Healthy Controls (HC), Group: II- Patients with T2DM without nephropathy (DM), Group: III- patients with T2DM with nephropathy (DM–CKD) in pre-dialysis stage (35) (Table 1). We have observed that increased level of inflammatory mediators such as TNF-α, hsCRP and MCP-1 may play independent as well as interdependent roles by influencing intracellular signalling which may contribute to hyperglycemia-mediated increase in inflammation and lead to development and progression of DN (35-39).

<table>
<thead>
<tr>
<th>Parameters</th>
<th>Group I (HC) (n = 50)</th>
<th>Group II (DM) (n = 50)</th>
<th>Group III (DM–CKD) (n = 50)</th>
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<tbody>
<tr>
<td>TNF-α (pg/mL)</td>
<td>14.5 ± 5.2 (13.1–16.1)</td>
<td>15.3 ± 3.7 (14.3–16.4)</td>
<td>20.6 ± 3.9 (19.5–21.8)</td>
</tr>
<tr>
<td>hsCRP (mg/L)</td>
<td>0.74 ± 0.46 (0.61–0.88)</td>
<td>3.6 ± 1.5a (3.3–4.1)</td>
<td>8.5 ± 1.7 (8.0–9.0)</td>
</tr>
<tr>
<td>uMCP-1 (pg/mg creatinine)</td>
<td>124.1 ± 46.6 (76.3–171.7)</td>
<td>278.5 ± 125.0 (153.1–400.8)</td>
<td>5632.7 ± 2275.8 (3351.5–8001.2)</td>
</tr>
</tbody>
</table>

Abbre: Group I- Healthy controls (HC), Group II- Diabetes mellitus (DM), Group III- Diabetes mellitus with CKD (DM–CKD). Tumor necrosis factor-alpha (TNF-α), High sensitive C-reactive protein (hsCRP), Urinary monocyte chemoattractant protein-1 (uMCP-1)
NF-κB1, which encodes for p105 subunit, that is ultimately processed to p50 subunit. TNF-α is a pro-inflammatory cytokine and both TNF-α and p85/p50 heterodimer (NF-kB) have been implicated in the pathogenesis of DN. We studied various genotypes (ins/ins, ins/del, del/del) of -94ins/del NF-kB1 gene to find their association in influencing the susceptibility of patients with DM to develop kidney disease. We studied NF-kB1 gene polymorphism in a total of 200 subjects which were divided in four groups of 50 subjects each: Group I-Healthy controls (HC), Group II- Diabetes mellitus (DM), Group III – DM with CKD (DM-CKD) and Group IV – Non-diabetic CKD (Non-DM-CKD). We observed that ins/del NF-kB1 genotype was present in highest number of subjects among all study groups except in patients with DM-CKD, where highest prevalence was of ins/ins genotype (Fig.1).

We also measured uMCP levels in these patients and found that these levels were significantly higher in patients of DM-CKD group as compared to HC and DM (p<0.001). These were also significantly higher than non-DM-CKD group (p<0.001). Further the patients with ins/ins NF-kB1 genotype had the highest level of uMCP suggesting the role of inflammatory pathway in pathogenesis of DN (Fig. 2).

![Graph showing distribution of genotypes](image_url)

**Fig.1:** Distribution of various genotypes of -94 insertion / deletion ATTG polymorphism of NF-kB1 gene in study groups
We found that the allele frequencies of -863C/A were 0.86/0.14 in HC group, 0.72/0.23 in patients with T2DM and 0.84/0.16 in DN, and that of -1031T/C were 0.89/0.11 in HC, 0.78/0.22 in DM, 0.87/0.13 in DM-CKD and 0.70/0.30 in Non DM-CKD.

Fig. 2: Urinary MCP-1 levels in relation to different NF-kB1 genotypes in various study groups

**Association of TNF-α promoter polymorphisms with plasma TNF-α levels and susceptibility to DN**

TNF-α is a pro-inflammatory cytokine which plays an important role in the pathogenesis of various inflammatory diseases including DN. Therefore, we evaluated the association of -863C/A (rs1800630) and -1031T/C (rs1799964) polymorphisms in the promoter region of TNF-α gene with plasma TNF-α levels among patients with T2DM with and without nephropathy.

We found that the allele frequencies of -863C/A were 0.86/0.14 in HC group, 0.72/0.23 in patients with T2DM and 0.84/0.16 in DN, and that of -1031T/C were 0.89/0.11 in HC, 0.78/0.22 in DM, 0.87/0.13 in DM-CKD and 0.70/0.30 in Non DM-CKD.

Table 2: Plasma TNF-α levels (pg/mL) in subjects with different TNF-α genotypes

<table>
<thead>
<tr>
<th>Groups</th>
<th>Gp I (HC) (n=100)</th>
<th>Gp II (DM) (n=100)</th>
<th>Gp III (DM-CKD) (n=100)</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Total</td>
<td>14.57±5.23</td>
<td>15.34±3.78</td>
<td>20.67±3.98,0.36</td>
<td></td>
</tr>
<tr>
<td>-863C→A</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>C/C</td>
<td>15.38±5.45</td>
<td>16.06±4.40</td>
<td>21.08±3.70</td>
<td>&lt;0.010</td>
</tr>
<tr>
<td>C/A</td>
<td>13.75±4.85</td>
<td>14.75±2.88</td>
<td>20.19±5.28</td>
<td>F-ratio=13.97 df=2</td>
</tr>
<tr>
<td>A/A</td>
<td>9.75±1.06</td>
<td>13.66±3.32</td>
<td>18.62±0.17</td>
<td></td>
</tr>
<tr>
<td>-1031T→C</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>T/T</td>
<td>14.31±5.16</td>
<td>15.18±3.91</td>
<td>19.44±3.60</td>
<td>0.104</td>
</tr>
<tr>
<td>Non T/T</td>
<td>15.37±5.62</td>
<td>16.76±1.94</td>
<td>23.29±3.52</td>
<td>F-ratio=1 df=4.71</td>
</tr>
</tbody>
</table>

Values are given as mean ± SD
Non T/T= T/T+C/C
Data are expressed as mean±SD. *p< 0.001 vs HC; †p<0.001 vs DM
Role of glutathione S-transferase M1 and T1 gene polymorphism in patients with DN

The glutathione S-transferases (GSTs) (EC 2.5.1.18) belong to a family of ubiquitous and multifunctional enzymes that work as one of the endogenous antioxidants through their ability to catalyze the conjugation of reduced GSH with electrophilic compounds and through their GSH peroxidase activity. Hence, reduced GST expression may result in diminished capacity of defense against OS. Interestingly, an earlier study has documented over-expression of GSTs in erythrocytes of CKD patients pointing to the fact that this group of enzymes might be involved in the pathogenesis of CKD (40). The mechanism by which GST polymorphism leads to CKD is not well-understood. In a previous study, Hayek et al (41) have shown that GSTT1 null genotype is associated with increase in markers of lipid peroxidation among diabetics.

In the last few years, there has been a surge of reports studying the association of DN with...
the genetic polymorphisms of GST. Although no association of GSTM1 deletion has been found with DN in Japanese T2DM patients (42), GSTT1 null genotype has been shown to be a risk factor for development of DN in the Chinese (43). In the Korean population (44), GSTM1 null genotype is found to be associated with nephropathy in T2DM patients. Ours is the first study regarding the association of GST polymorphism with DN in Indian population.

We investigated the role of the GST polymorphisms in determining variation in susceptibility of individuals to CKD and to pinpoint the probable underlying mechanism (45, 46). A total of 200 subjects were enrolled under four study groups, viz., (i) Healthy subjects, n = 50; (ii) Patients of T2DM for at least 10 years without any microalbuminuria or overt proteinuria, n = 50; (iii) Patients of T2DM with nephropathy (DM-CKD), characterized by the presence of microalbuminuria or overt proteinuria, n = 50; (iv) Patients with nondiabetic CKD (NDM-CKD) having evidence of overt proteinuria and/or deranged renal function for more than 3 months in the absence of DM and any systemic or local infection, n = 50. We found that GSTM1 and GSTT1 deletions singly or together were associated with lower GST levels and higher OS in both diabetic and nondiabetic CKD. Interestingly GSTT1 deletion appears to be associated with both diabetic and nondiabetic CKD irrespective of the GSTM1 status.

Role of GSTM1 and GSTT1 genotypes in the development of OS in patients with DN

In another study, we investigated whether the deletion of GSTM1 and GSTT1 genes are associated with higher OS in DN patients and the possible role of these polymorphisms in the development of DN (47, 48). We recruited 60 patients having T2DM for more than 5 years with albuminuria >300 mg/day and having evidence of diabetic retinopathy. In all these patients genotypic analysis was done by using multiplex polymerase chain reaction (PCR). Various OS parameters which were studied included GST activity, MDA and GSH. The prevalence of different genetic polymorphisms of GST in patients with DN was: GSTM1 positive - 32 (53.4%), GSTM1 null – 28 (46.6%), GSTT1 positive – 27 (45%), GSTT1 null – 33 (55%), Both null – 18 (30%) and Both positive – 17 (28.3%).

Further, our study showed that double deletion involving GSTT1 and GSTM1 may result in decreased GST levels, leading to increased OS which may result in greater risk for development of DN. The levels of different OS parameters in various GST genotypes are given in Table 4.

<table>
<thead>
<tr>
<th>GSTM1 null(n = 28)</th>
<th>0.95 ± 0.16</th>
<th>5.29 ± 0.89</th>
<th>142.1 ± 26</th>
</tr>
</thead>
<tbody>
<tr>
<td>GSTT1 null(n = 33)</td>
<td>0.98 ± 0.17</td>
<td>5.17 ± 1.14</td>
<td>143.4 ± 33.3</td>
</tr>
<tr>
<td>Both null(n = 18)</td>
<td>0.87 ± 0.11a,b</td>
<td>5.84 ± 0.6a,b,c</td>
<td>130.6 ± 34a</td>
</tr>
<tr>
<td>Both positive(n = 17)</td>
<td>1.27 ± 0.12</td>
<td>4.28 ± 1.19</td>
<td>173.9 ± 24.5</td>
</tr>
</tbody>
</table>

*aSignificantly different from both positive group at p<0.05,  
'bSignificantly different from GSTT1 null group at p<0.05,  
'cSignificantly different from GSTM1 null group at p<0.05  
(ANOVA with Tukey SD as post-hoc test)
**GST levels in patients with T2DM with and without nephropathy**

Hyperglycemia induced OS is implicated as a contributor to the onset and progression of T2DM and its complications like DN. GST is primarily involved in the neutralization of Reactive Oxygen Species (ROS) by enzymatic conjugation with the scavenger peptide GSH (49-51). In another study, we evaluated the role of GST along with OS markers and their correlation in patients with T2DM with and without nephropathy. We have recruited 300 study subjects divided into three groups of 100 each: HC, DM and DM-CKD. Plasma GST, MDA, reduced GSH levels and FRAP were estimated spectrophotometrically.

GST levels were found to be raised in the patient groups in comparison to HC, however, the highest significant levels were seen in T2DM as compared to DM-CKD (p<0.05). Other oxidant and antioxidant markers are shown in Table 5. The antioxidant parameters including GSH and FRAP were significantly lower in DN as compared to T2DM and HC (p<0.05). Higher lipid peroxidation was seen in patients of DM-CKD as MDA levels were significantly raised when compared to T2DM and HC (p<0.05). We found negative correlation between HbA1c and GSH (r= -0.942, p<0.01) and FRAP (r= -0.854, p<0.01). A positive association was observed between HbA1c and GST (r= 0.606, p<0.01) and MDA (r= 0.839, p<0.01); however, a significant negative correlation between GST activity and GSH levels (r= -0.530, p<0.01) and FRAP (r= -0.294, p<0.01) was observed. GST activity was positively correlated with MDA levels (r= 0.253, p<0.01).

**Table 5: Plasma levels of oxidant-antioxidant markers in various study groups**

<table>
<thead>
<tr>
<th>Parameters</th>
<th>Group I (HC) (n=100)</th>
<th>Group II (DM) (n=100)</th>
<th>Group III (DM-CKD) (n=100)</th>
</tr>
</thead>
<tbody>
<tr>
<td>GSH (mg/g Hb)</td>
<td>3.2±0.2 (3.0-3.9)</td>
<td>1.7±0.2&lt;sup&gt;a,c&lt;/sup&gt; (1.0-1.9)</td>
<td>0.85±0.1&lt;sup&gt;a,b&lt;/sup&gt; (0.12-1.1)</td>
</tr>
<tr>
<td>GST (nmol/min )</td>
<td>6.3±0.6 (4.1-7.8)</td>
<td>8.6±0.5&lt;sup&gt;a,c&lt;/sup&gt; (7.6-10.2)</td>
<td>7.5±0.3&lt;sup&gt;a,b&lt;/sup&gt; (6.25-8.1)</td>
</tr>
<tr>
<td>MDA (nmol/mL)</td>
<td>1.5±0.3 (0.9-2.2)</td>
<td>2.5±0.3&lt;sup&gt;a,c&lt;/sup&gt; (2.0-3.1)</td>
<td>5.2±0.5&lt;sup&gt;a,b&lt;/sup&gt; (4.3-6.8)</td>
</tr>
<tr>
<td>FRAP (µmol/L)</td>
<td>542.9±54.0 (425-660)</td>
<td>404.8±48.4&lt;sup&gt;a,c&lt;/sup&gt; (300.8-494.9)</td>
<td>163±40.2&lt;sup&gt;a,b&lt;/sup&gt; (95.5-335)</td>
</tr>
</tbody>
</table>

Data are expressed as mean±SD. Group I- Healthy control (HC); Group II- Type 2 diabetes mellitus without nephropathy (DM) Group III- Diabetes mellitus with CKD (DM-CKD).

<sup>a</sup>significantly different from healthy control at p<0.05,

<sup>b</sup>significantly different from diabetic patients without nephropathy at p<0.05,

<sup>c</sup>significantly different from diabetic patients with nephropathy at p<0.05.

**D. Genes Involved in Cytoprotective Pathway**

NAD(P)H quinone oxidoreductase 1 (NQO1) plays a prominent role in maintaining cellular homeostasis and is an essential component of the antioxidant defense system. It catalyzes metabolic detoxification of quinines and protects cells against chemical induced OS. Expression of NQO1 increases in response to oxidant and electrophilic radicals to counteract OS, and in fact, it is a kind of cytoprotective defense mechanism for cells.
Table 6. NQO1 levels in plasma (ng/mL) in context of different NQO1 genotypes

<table>
<thead>
<tr>
<th>Groups</th>
<th>Group I (HC) (n=200)</th>
<th>Group II (DM) (n=200)</th>
<th>Group III (DN) (n=200)</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>NQO1*1/*1</td>
<td>4.16±0.47</td>
<td>31.36±1.19</td>
<td>15.51±0.78</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>NQO1*1/*2</td>
<td>1.9±0.46&lt;sup&gt;a&lt;/sup&gt;</td>
<td>25.99±0.81&lt;sup&gt;a&lt;/sup&gt;</td>
<td>11.06±0.67&lt;sup&gt;a&lt;/sup&gt;</td>
<td></td>
</tr>
<tr>
<td>NQO1*2/*2</td>
<td>0.62±0.06&lt;sup&gt;a,b&lt;/sup&gt;</td>
<td>20.32±1.28&lt;sup&gt;a,b&lt;/sup&gt;</td>
<td>7.43±0.85&lt;sup&gt;a,b&lt;/sup&gt;</td>
<td></td>
</tr>
</tbody>
</table>

Plasma NQO1 levels are given as mean±SD. Group I- Healthy Control (HC), Group II- Type 2 Diabetes Mellitus without complications (DM), Group III- Diabetic Nephropathy (DN).<sup>a</sup>significantly different from homozygous NQO1*1/*1 in study groups at p<0.05<sup>b</sup>significantly different from heterozygous NQO1*1/*2 in study groups at p<0.05

Among DN and T2DM patients, the OR for the development of DN was 1.72-fold higher in T2DM patients carrying the NQO1*2/*2 genotype than in those carrying NQO1*1/*1 or NQO1*1/*2 genotypes (95% CI=1.133 to 2.600). A significant association was observed for NQO1*2 polymorphism in patients of T2DM when compared to HC (OR=6.638, 95% CI=1.427–30.876, p=0.016). NQO1*2 polymorphism was shown to be associated with an increased risk of DN in comparison to HC (OR=22.00, 95% CI=5.075–95.376, p=0.000). Therefore, it may be concluded that NQO1*2 allele may increase the risk for developing DN in T2DM patients as well as HC.

E. Genes Involved in Nitric Oxide (NO) Pathway

NO is a major regulator of renal hemodynamics, its production being catalysed by endothelial nitric oxide synthase (eNOS). Reduction in the generation of NO acts as a deteriorating factor for progressive renal disease. Polymorphisms in the eNOS gene may alter its expression, thus affecting the production of NO. Familial clustering DN points to a role of genetic factors in the pathogenesis of renal disease.
Hey et al published a meta-analysis of 24 studies and analysed the polymorphisms of eNOS genes (4b/a,G894T and T786C) associated with DN (55). It was found that 4b/a and G894T polymorphisms in the eNOS gene were associated with susceptibility to DN in Asian populations, but not in Caucasian populations. A meta-analysis of 8 studies performed by Zhou et al evaluated the association of G894T gene polymorphism alone with DN susceptibility (56). These studies included 850 cases and 1254 controls. In the Asian population, the average frequency of T-allele was 19.19% in DN patients and 8.68% in controls. A significant association was observed between the presence of T-allele and DN risk in the overall population.

We studied the eNOS G894T polymorphism in patients of diabetes with and without nephropathy and measured the serum NO levels in these patients and compared them in the HCs. It was found that patients with DN add lowest levels of NO (22.02±16.18 µM) as compared to diabetics without nephropathy (63.86 ± 29.49 µM) and HC (38.42 ± 13.71 µM) (p< 0.001). Further a positive association was observed between eNOS G894T polymorphism as the frequency of TT genotype as well as that of mutant T allele was increased in patients with DN as compared to diabetics without nephropathy and HCs (57).

F. Genes Involved in Advanced Glycation End-products (AGE) and its Receptor (RAGE) Interaction

Association of RAGE gene polymorphism with vascular complications in patients with T2DM

Hyperglycemia associated with DM stimulates non-enzymatic glycation and oxidation of proteins and lipids leading to enhanced formation of AGEs. There is growing evidence that production and accumulation of AGEs is involved in the initiation and development of micro- and macrovascular complications observed in DM (58, 59). AGEs bind to specific RAGE which is expressed in many of the cell types such as endothelial cells, monocytes and lymphocytes, including β cells of pancreas (60-62). The gene for RAGE is located on the chromosome 6p21.3 near the HLA locus, and at least 30 polymorphisms have been identified of which 9 are in promoter region, 11 in exon region and 10 in intron region.

Three polymorphisms of RAGE gene namely -374T/A, -429T/C and Gly82Ser have been widely studied with regard to the development of diabetic complications in different populations all over the world. We investigated the association of -374T/A, -429T/C and Gly82Ser RAGE gene polymorphisms and their haplotypes with vascular complications in T2DM patients which may help in identifying DM patients predisposed to possible micro- and macrovascular complications as a result of their genetic makeup (63). A total of 427 patients of T2DM with disease duration > 5 years were enrolled in this study. These patients were divided into three groups. The first group referred to as 'DM' comprised of 140 T2DM subjects without any vascular complications. The second group referred to as 'DM-micro' consisted of 152 T2DM subjects with microvascular complications (retinopathy and nephropathy). The third group referred to as 'DM-macro' consisted of 135 T2DM subjects with macrovascular complications. In addition, 176 HCs were enrolled in the study.

We found that -429T/C and Gly82Ser RAGE polymorphisms were significantly associated with the development of macrovascular and microvascular complications respectively in T2DM subjects while -374A allele showed reduced risk towards the development of macrovascular complications. Further, -429T/C, -374T/A and Gly82Ser haplotype analysis revealed association of CTG haplotype with development of macrovascular complications while haplotype TAG was observed to be significantly protective towards development of macrovascular complications in
T2DM subjects (OR = 0.617, p = 0.0202).

G. Gene Expression Studies

Role of AGE-RAGE expression in diabetic vascular complications

Interaction of AGE with its receptor RAGE transduces multiple signals such as NAD(P)H oxidase, Mitogen-Activated Protein (MAP) kinases, extracellular signal regulated kinases, GTPase, etc. (64, 65). Activation of NAD(P)H oxidase causes enhanced Reactive Oxygen Species (ROS) generation which may lead to peroxidation and glycoxidation reactions that result in Protein Carbonyl (PCO) formation, Advanced Oxidation Protein Products (AOPP) generation and lipid peroxidation. These OS markers have been shown to be enhanced significantly in diabetic patients (66, 67). On the other hand, activation of kinases and GTPases causes activation of nuclear transcription factor including NF-kB which transcribes its target genes such as Vascular Cell Adhesion Molecule – 1 (VCAM-1), E-selectin and pro-inflammatory cytokines.

AGE-RAGE interaction is one of the mediators of vascular complications in DM; however, factors that possibly induce exaggerated AGE-RAGE interaction are not well known. RAGE is usually expressed at low levels in adults. Enhanced AGE-RAGE interaction possibly requires increased expression of RAGE. In various diseased states such as CVD, diabetes, inflammation, etc., there is higher expression of RAGE; however, conditions and factors that may induce RAGE expression particularly in T2DM have not been elucidated.

Since AGE formation is an integral phenomenon in T2DM, we investigated the dependence of RAGE expression on circulating AGE level and have examined the outcome of AGE-RAGE interaction by measurement of OS status in those patients (68). We recruited 75 patients of T2DM with disease duration > 5 years for this study. These patients were divided into three groups. The first group referred to as ‘DM’ comprised of 25 T2DM subjects without any vascular complications. The second group referred to as ‘DM-micro’ consisted of 25 T2DM subjects with microvascular complications (retinopathy and nephropathy). The third group referred to as ‘DM-macro’ consisted of 25 T2DM subjects with macrovascular complications. In addition, 25 HCs were also enrolled in the study by voluntary participation.

We observed that serum AGEs levels were significantly higher in diabetic patients having vascular complications as compared to T2DM without complications (p < 0.01) (Table 7). RAGE m-RNA expression level in PBMCs assayed by quantitative real time PCR was four times higher in diabetic subjects without vascular complications while DM patients having microvascular or macrovascular complications showed 12 fold and 8 fold higher RAGE m-RNA expression, respectively, compared to HCs. Further, circulating AGEs levels showed significant positive correlation with RAGE m-RNA expression and OS markers (Fig.3).

H. Epigenetic Studies

Epigenetics is the study of inherital changes in phenotype or gene expression caused by mechanisms other than changes in the underlying DNA sequence; therefore, non-genetic factors cause the organism's genes to behave differently. There is some evidence to suggest the role of epigenetic factors in the complex interplay between genes and environment. Epigenetic mechanisms include DNA methylation, histone modification and microRNAs. Few studies have suggested that hyperglycemia may induce epigenetic changes of pro-inflammatory genes, which subsequently regulate gene expression and thereby the development of vascular complications (69, 70); however, improved glycemic control for 3 – 5 years at a later stage in diabetic patients did not reduce the risk of macrovascular complications.
Table 7. Serum AGE levels and oxidative stress markers in diabetic patients and healthy subjects

<table>
<thead>
<tr>
<th>Parameters</th>
<th>DM</th>
<th>DM-micro</th>
<th>DM-macro</th>
<th>Healthy subjects</th>
</tr>
</thead>
<tbody>
<tr>
<td>AGE ELISA (µg/mL)</td>
<td>1.4±0.54</td>
<td>3.4±0.95a</td>
<td>2.32±0.88a</td>
<td>1.12±0.38</td>
</tr>
<tr>
<td>AGE-F (AU)</td>
<td>1.97±0.43</td>
<td>2.68±0.39a</td>
<td>2.50±0.32a</td>
<td>1.87±0.29</td>
</tr>
<tr>
<td>MDA (nmol/mL)</td>
<td>0.43±0.15</td>
<td>0.76±0.22a</td>
<td>0.81±0.37a</td>
<td>0.26±0.06</td>
</tr>
<tr>
<td>PCO (nmol/mg protein)</td>
<td>1.6±0.43</td>
<td>2.8±1.0a</td>
<td>3.0±0.72a</td>
<td>1.4±0.43</td>
</tr>
<tr>
<td>AOPP (µmol/L of chloramines T-equivalent)</td>
<td>103±27.2</td>
<td>163±42.2a</td>
<td>206±54.2a</td>
<td>85±21.9</td>
</tr>
</tbody>
</table>

Data are presented as mean ± SD.

Comparison between the groups was performed with one-way ANOVA and followed by post hoc Tukey's analysis. *p < 0.05 compared with controls and DM group.

Advanced glycation end-product (AGE), Malonyldialdehyde (MDA), Protein carbonyl (PCO), Advanced oxidation protein product (AOPP), Diabetes mellitus (DM), Diabetes mellitus with microvascular complications (DM-micro), Diabetes mellitus with macrovascular complications (DM-macro).

Fig. 3: Circulating AGEs shows significant positive correlation with RAGE m-RNA expression among diabetic subjects. (A) Relationship between circulating AGEs and RAGE m-RNA expression among diabetic subjects. (B) Relationship between fluorescent AGEs and RAGE m-RNA expression among diabetic subjects. Correlation analysis was performed using Pearson's coefficient.
significant association of -429T/C and Gly82Ser RAGE polymorphisms with the development of macrovascular and microvascular complications, respectively in T2DM subjects while -374A allele showed reduced risk towards the development of macrovascular complications. Further, we have observed that AGE-mediated exacerbation of RAGE expression may contribute to OS generation that plays a key role in pathogenesis of various vascular complications in DM.

To conclude, polymorphisms of various genes involved in RAAS, inflammatory, oxidant stress, cytoprotective and nitrous oxide pathways may influence the levels of various cytokines and intracellular signaling mechanisms, thereby influencing the susceptibility of patients with DM for development of kidney disease. In addition, raised AGE and AOPP levels and polymorphisms of RAGE and enhanced RAGE mRNA expression may adversely influence final common pathway through oxidant stress mechanisms, and thereby determine the development of microvascular and/or macrovascular complications in patients with DM.

Key Findings and Summary

There is enough evidence from our studies to support the hypothesis that genetic factors play a crucial role in determining susceptibility of patients with DM for development of nephropathy. We have found that D allele of ACE gene acts as risk factor for the development of nephropathy in patients with T2DM as well as for nondiabetic CKD; whereas I allele of ACE gene was protective in nature. Further, whether ACE gene polymorphism can modulate the degree of beneficial reno-protective effect following ACE inhibitor therapy is currently under investigation in our laboratory. We have found that OS plays a significant role in the development of DN, and that GSTT1 and/or GSTM1 null genotypes are associated with higher OS in patients with DN. In addition, we also found that increased levels of inflammatory mediators i.e. TNF-α, hsCRP and uMCP-1 play an independent as well as interdependent roles, via several signaling pathways contributing to hyperglycemia-mediated increase in OS. We have shown that genetic polymorphism of NF-kB gene and TNF-α gene plays a significant role in determining serum level of various inflammatory markers and OS parameters. An increase in OS may further amplify inflammation, thus setting up a vicious cycle. Therefore, inflammation interlinked with OS may be major mechanisms in the pathogenesis and progression of nephropathy in susceptible diabetic patients. We found significant association of -429T/C and Gly82Ser RAGE polymorphisms with the development of macrovascular and microvascular complications, respectively in T2DM subjects while -374A allele showed reduced risk towards the development of macrovascular complications. Further, we have observed that AGE-mediated exacerbation of RAGE expression may contribute to OS generation that plays a key role in pathogenesis of various vascular complications in DM.

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Higher plasma levels of advanced glycation end products are associated with incident cardiovascular disease and all-cause mortality in type 1 diabetes. *Diabetes Care* **34**:442–447.


Essential Skills in Postgraduate Medical Curriculum of Community Medicine

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ABSTRACT

**Introduction:** Community-based education has been considered a suitable approach for health promotion and for requisite skill development regarding primary health care. In the current perspective, public health training and research, being two important aspects require immediate attention.

**Objective:** To assess the skills of Postgraduate Students in the Department of Community Medicine in four Medical Colleges of Delhi.

**Materials and Methods:** It was a cross-sectional study conducted among 70 Postgraduate Medical Students of 4 Medical Colleges in Delhi. The data were collected through a self administered, pre-tested questionnaire containing items assessing socio-demographic profile and skills essential for Postgraduate Students of Community Medicine.

**Results:** There were 58.6% male and 29% female students. A large proportion of participants were having age range between 25-29 years. Ability 'to resolve conflict among the nurse at Primary Health Centre (PHC)', 'generate community participation', 'making thick and thin smear in case of fever', 'making a chart showing month-wise distribution of CuT', and 'calculating Chi-square of data', were found to significantly higher in 2nd and 3rd year PG students than first year PG students (p<0.01). Only 27.1% of students felt that they could test water sample for microbiological aspects while only 47.1% said that they could examine an industrial worker for pre-placement examination.

**Conclusions:** PG students assessed themselves to possess necessary skills on communication, counselling and health education. However, many students lacked skills pertaining to occupational health and epidemiology.

**Keywords:** Competency in community medicine, public health, epidemiological skills, communication skills.

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GOLDEN JUBILEE COMMEMORATION AWARD LECTURE delivered during NAMSCON 2016 at the All India Institute of Medical Sciences, Raipur.
Introduction

Based on the principles outlined at Alma-Ata in 1978, there is an urgent call for revitalizing primary health care. India is not only committed to strengthening primary care but is already well aware of the means to achieving this goal (1). The National Rural Health Mission (NRHM) is a great testament to the determination of the Indian government to deliver universal primary health care and it has had impact also (2). However, there are a number of operational issues that need to be addressed to ensure that primary care is delivered effectively. And, some of the major problems in implementation and practice of primary health care relate to training and capacity building of health service providers in foreseeable future (3). It is suggested that human resources with requisite knowledge, skills, attitudes, values and responsiveness to people's health needs and health promotion are needed.

Community-based education has been considered a suitable approach for health promotion and for requisite skill development regarding primary health care (4, 5). In the current perspective, public health training and public health research, being two important aspects require immediate attention. The Task Force on Medical Education for the NRHM and National Health Policy 2002 have recommended increase in postgraduate seats in the discipline of Community Health/Public Health/Preventive, Social Medicine and Family Medicine (3). Thus, it is evident that in current scenario, public health specialists are the need of the hour in community health care.

The objectives and goals of Postgraduate (PG) Medical Education in Community Medicine are to produce competent specialists to manage the teaching departments in the Medical Colleges, or to manage health services and national health programs at various levels or to conduct biomedical research in the discipline of Community Medicine (6). During training, PG are expected to acquire a substantial knowledge and necessary skills in: concepts of health and illness and their determinants, methods in community health, health services organizations, community health programs and communication and advocacy (7).

There are regional variations in the way PG training in India is imparted by various institutions. There is no well defined curriculum worked out, if at all it is there, the methods of acquiring the desired competencies are vague or not at all stated (6). Though Medical Council of India has reiterated competency-based PG training in community health care, yet, detailed structured curriculum in PG training in Community Medicine is being followed in handful institutions only (8).

Considering the fact acquiring relevant public health skills during PG training is pertinent, the present study was undertaken to assess the skills of PG students in department of Community Medicine in four Medical Colleges of Delhi. This would also enable us to know the variations in PG training and evaluate whether the medical pedagogic approach is in line with objectives of such training.

Materials and Methods

Study Setting and Study Participants

It was a cross-sectional study conducted among PG Medical Students (Doctor of Medicine: MD) of 4 Medical Colleges in Delhi. Every year, 8-9 students are admitted in each of the Medical Colleges; therefore, there are around 24-27 students in each of the Department of Community Medicine of a Medical College, studying at any given time. The entire course of MD is of 3 years. The PG Students are posted for practical training in primary health care in Department of Community Medicine from first year onwards.
**Description of Curriculum**

As per MCI 2000 regulations, the major components of PG Curriculum in Community Medicine constitutes:

1. Theoretical knowledge
2. Practical and clinical skills (Managerial skills)
3. Thesis skills
4. Attitudes including communication skills
5. Training in research methodology.

In order to achieve this practical training in postgraduation in Medical Colleges of Delhi is divided as follows:

The PG students are posted on rotational basis in the field practice areas of their respective Departments in different Medical Colleges for their practical training. The field areas usually cover the urban slums or resettlement colonies and rural areas include villages. The posting is of 3-6 months duration in each area. During this posting, PG students are expected to know the community, establish rapport with the community and provide essential primary care to the community. They study the community and learn to make community diagnosis. Further, they learn the implementation of national health programs at field level. They are also expected to learn management of human resources. Sometimes, their research work is also carried out in these areas.

PG students are also posted in Family Health Care program (FHCP) for undergraduates which is covered in 4th and 5th semester and epidemiology and biostatistics which are covered in 6th and 7th semester. Besides, some extra-mural visits are arranged during this posting that includes visit to a health centre, Anganwadi, Medical Records Department, Nursing College for injection techniques, MCH centre, Institute of Physically Handicapped (IPH), and Centre for Environmental and Occupational Health (CEOH) for Biomedical Waste Management training. In addition to this, health communication classes are also held where in students learn the skills of imparting health education in a community.

PG students are also scheduled to present journal articles and take seminars on topics of public health importance. It is mandatory to do research work towards a thesis for award of the PG degree in their third year of postgraduation, they are also posted in various public health institutes and agencies like National Centre for Disease Control (NCDC), National Institute of Health and Family Welfare (NIHFW), Municipal Corporation of Delhi (MCD), New Delhi Municipal Corporation (NDMC), etc. They are also posted in the Department of Microbiology of their respective colleges for learning organisms of public health importance. Additionally, the students participate in seminars, symposia, conferences and workshops for enhancing their knowledge about the advances in public health. They are encouraged to present their research work in these fora.

The students are finally evaluated at the end of third year. There are 4 theory papers followed by 2-day practical exam that includes long case, two short cases, epidemiological and statistical exercises, spots and viva voce. Some colleges also hold microbiology practicals.

**Sample Size and Sampling**

There are about 8-9 PG seats for the subject in the studied Medical Colleges of Delhi. All the PG Students during the study period were included in the study. There were total of 96 PG seats in three years in the four Medical Colleges (24 in each college). Out of the 96, only 90 were filled and only 70 returned the completed questionnaires. Hence, they were included for analysis.

**Study Tool**

The study tool was developed after extensive review of curriculum of
Postgraduation in Community Medicine in various institutes. The questionnaire so developed was self-administered, pre-tested and consisted of two parts. The first part contained items to gather socio-demographic information of the participants. The second part consisted of various questions to assess public health skills. To identify skills needed for PG students, emails to 50 faculty of Community Medicine were sent and their responses were listed. Questions were framed covering all skills suggested by faculty members. The skills that were assessed covered following areas: communication with the community, school health, human resource management, social mobilization, laboratory services, management of public health problem, family planning counselling, graphic representation of data, immunization, nutrition, water testing and disinfection, occupational health, health education, investigation of an epidemic, statistical tests, inventory management, evaluation of national health program, qualitative research, and surveillance.

**Methodology**

All the PG students of four medical colleges were contacted and the objective of study was explained to them. Informed written consent was obtained from them and then the questionnaires were given to them. The students were asked to return the filled questionnaire in 3 days. Those who failed to return were contacted again. Those students who failed to submit the filled questionnaire even after 2 remainders were excluded from the study.

**Ethical Considerations**

The privacy and confidentiality of data was ensured. The ethical approval of the research was obtained from the Institutional Ethics Committee (IEC) of the research institute.

**Statistical Analysis**

The obtained data were collected, entered and analyzed in SPSS (version 17.0). Data were expressed in terms of percentages. Differences between the proportions were observed by Chi-square test and p < 0.05 was considered as significant. Difference between independent groups for continuous variables was assessed using student t test (2 groups) and One-way ANOVA (for more than 2 groups).

**Results**

Out of the total 70 respondents, 58.6% were male and 29% were females. Large proportion of participants was between the age of 25-29 years. More than half of the participants had their undergraduation from a private college (51.4%). Majority of the students belonged to general category (n=49; 70%). There were 29 students who were in first year of postgraduation, 18 were in 2nd year while 23 students were studying in 3rd year. While 34.3% students had graduate mothers, half of the PG students had Fathers with professional education. Majority of students said that their mothers were housewives (87.1%) while half had fathers engaged in semi-professional occupation (Table 1).

There were 22 competencies that were self-assessed by the students. Most of the competencies did not show year of PG wise difference. However, ability 'to resolve conflict among the nurse at Primary Health Centre (PHC)', 'generate community participation', 'making thick and thin smear in case of fever', 'making a chart showing month-wise distribution of CuT', and 'calculating Chi-square of data', was found to be significantly higher in 2nd and 3rd year PG students than first year PG students (p<0.01). Only 27.1% of students felt that they could test water sample for microbiological aspects while only 47.1% said that they could examine an industrial worker for pre-placement examination. Ability to evaluative DOTS or any national health program and manage inventory of health centre was possessed by 48.6% and 54.3%, respectively (Table 2).
Table 1: Socio-demographic Characteristics of Students (n=70)

<table>
<thead>
<tr>
<th>Variables</th>
<th>Frequency (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Gender</strong></td>
<td></td>
</tr>
<tr>
<td>- Male</td>
<td>41 (58.6)</td>
</tr>
<tr>
<td>- Female</td>
<td>29 (41.4)</td>
</tr>
<tr>
<td><strong>Age</strong></td>
<td></td>
</tr>
<tr>
<td>- 20-24 years</td>
<td>4 (5.7)</td>
</tr>
<tr>
<td>- 25-29 years</td>
<td>61 (87.2)</td>
</tr>
<tr>
<td>- 30-34 years</td>
<td>5 (7.1)</td>
</tr>
<tr>
<td><strong>Type of medical college (where MBBS done)</strong></td>
<td></td>
</tr>
<tr>
<td>- Government</td>
<td>34 (48.6)</td>
</tr>
<tr>
<td>- Private</td>
<td>36 (51.4)</td>
</tr>
<tr>
<td><strong>Category</strong></td>
<td></td>
</tr>
<tr>
<td>- General</td>
<td>49 (70.0)</td>
</tr>
<tr>
<td>- Scheduled Caste (SC)</td>
<td>9 (12.9)</td>
</tr>
<tr>
<td>- Scheduled Tribe (ST)</td>
<td>5 (7.1)</td>
</tr>
<tr>
<td>- Other Backward Class (OBC)</td>
<td>7 (10.0)</td>
</tr>
<tr>
<td><strong>Year of Postgraduation</strong></td>
<td></td>
</tr>
<tr>
<td>- 1\textsuperscript{st} year</td>
<td>29 (41.4)</td>
</tr>
<tr>
<td>- 2\textsuperscript{nd} year</td>
<td>18 (25.7)</td>
</tr>
<tr>
<td>- 3\textsuperscript{rd} year</td>
<td>23 (32.9)</td>
</tr>
<tr>
<td><strong>Mother’s education</strong></td>
<td></td>
</tr>
<tr>
<td>- Professional</td>
<td>16 (22.8)</td>
</tr>
<tr>
<td>- Graduate</td>
<td>24 (34.3)</td>
</tr>
<tr>
<td>- 12\textsuperscript{th} standard</td>
<td>11 (15.8)</td>
</tr>
<tr>
<td>- 10\textsuperscript{th} standard</td>
<td>11 (15.8)</td>
</tr>
<tr>
<td>- Middle school</td>
<td>2 (2.8)</td>
</tr>
<tr>
<td>- Primary school</td>
<td>1 (1.4)</td>
</tr>
<tr>
<td>- Illiterate</td>
<td>5 (7.1)</td>
</tr>
<tr>
<td><strong>Father’s education</strong></td>
<td></td>
</tr>
<tr>
<td>- Professional</td>
<td>35 (50)</td>
</tr>
<tr>
<td>- Graduate</td>
<td>26 (37.1)</td>
</tr>
<tr>
<td>- 12\textsuperscript{th} standard</td>
<td>4 (5.7)</td>
</tr>
<tr>
<td>- 10\textsuperscript{th} standard</td>
<td>3 (4.3)</td>
</tr>
<tr>
<td>- Primary school</td>
<td>1 (1.4)</td>
</tr>
<tr>
<td>- Illiterate</td>
<td>1 (1.4)</td>
</tr>
<tr>
<td><strong>Mother’s occupation</strong></td>
<td></td>
</tr>
<tr>
<td>- Professional</td>
<td>2 (2.9)</td>
</tr>
<tr>
<td>- Semi-professional</td>
<td>7 (10.0)</td>
</tr>
<tr>
<td>- Housewives</td>
<td>61 (87.1)</td>
</tr>
<tr>
<td><strong>Father’s occupation</strong></td>
<td></td>
</tr>
<tr>
<td>- Professional</td>
<td>11 (15.8)</td>
</tr>
<tr>
<td>- Semi-professional</td>
<td>35 (50)</td>
</tr>
<tr>
<td>- Clerical, shop owners, etc.</td>
<td>12 (17.1)</td>
</tr>
<tr>
<td>- Skilled worker</td>
<td>2 (2.9)</td>
</tr>
<tr>
<td>- Unskilled worker</td>
<td>1 (1.4)</td>
</tr>
<tr>
<td>- Unemployed</td>
<td>9 (12.9)</td>
</tr>
</tbody>
</table>
Table 2: Skill Assessment of Postgraduate Students

<table>
<thead>
<tr>
<th>Skill</th>
<th>Year of Postgraduation (n, %)</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>1&lt;sup&gt;st&lt;/sup&gt; Year</td>
<td>2&lt;sup&gt;nd&lt;/sup&gt; Year</td>
</tr>
<tr>
<td>Communicate with a senior family member at home and in field to get the information about socio-demography</td>
<td>26 (89.7)</td>
<td>15 (83.3)</td>
</tr>
<tr>
<td>Plan and carrying out school health program in a school</td>
<td>16 (55.2)</td>
<td>11 (61.1)</td>
</tr>
<tr>
<td>Resolve conflict between two nurses working under you at Primary Health Centre (PHC)</td>
<td>14 (48.3)</td>
<td>16 (88.9)</td>
</tr>
<tr>
<td>Generate community participation for clean surrounding of their houses</td>
<td>15 (51.7)</td>
<td>15 (83.3)</td>
</tr>
<tr>
<td>Negotiate with local NGO’s for decreasing female feticide/ANC/Institutional delivery</td>
<td>16 (55.2)</td>
<td>11 (61.1)</td>
</tr>
<tr>
<td>Make thick and thin smear at home of patient with fever</td>
<td>10 (34.5)</td>
<td>13 (72.2)</td>
</tr>
<tr>
<td>Categorize TB patients based on the sputum exam and history at PHC</td>
<td>26 (89.7)</td>
<td>17 (94.4)</td>
</tr>
<tr>
<td>Motivate a couple to use contraceptives of their choice</td>
<td>27 (93.1)</td>
<td>18 (100)</td>
</tr>
<tr>
<td>Make chart showing month-wise distribution of Copper T users</td>
<td>5 (17.2)</td>
<td>16 (88.9)</td>
</tr>
<tr>
<td>Plot weight of a &lt;5 child on “Road to health” card</td>
<td>25 (86.2)</td>
<td>17 (94.4)</td>
</tr>
<tr>
<td>Give BCG vaccine to an infant</td>
<td>28 (96.6)</td>
<td>17 (94.4)</td>
</tr>
<tr>
<td>Plan a diet for a child suffering from grade 2 PEM</td>
<td>17 (58.6)</td>
<td>14 (77.8)</td>
</tr>
<tr>
<td>Chlorinate one bucket of water at home</td>
<td>17 (58.6)</td>
<td>10 (55.6)</td>
</tr>
<tr>
<td>Examine thoroughly an industrial worker for pre-placement examination</td>
<td>11 (37.9)</td>
<td>9 (50)</td>
</tr>
<tr>
<td>Deliver a health talk on diarrhoea prevention and control at health centre</td>
<td>27 (93.1)</td>
<td>17 (94.4)</td>
</tr>
<tr>
<td>Investigate an epidemic of food poisoning in a village/hostel</td>
<td>14 (48.3)</td>
<td>12 (66.7)</td>
</tr>
<tr>
<td>Calculate Chi-square value for a given data</td>
<td>13 (44.8)</td>
<td>13 (72.2)</td>
</tr>
<tr>
<td>Manage the inventory/drug store at the health centre</td>
<td>15 (51.7)</td>
<td>7 (38.9)</td>
</tr>
<tr>
<td>Evaluate a DOTS/Any other program in your district/state</td>
<td>14 (48.3)</td>
<td>6 (33.3)</td>
</tr>
<tr>
<td>Hold a focus group discussion for evaluating barriers in early initiation of breastfeeding</td>
<td>24 (82.8)</td>
<td>16 (88.9)</td>
</tr>
<tr>
<td>Test the water sample for micro-biological aspects</td>
<td>8 (27.6)</td>
<td>4 (22.2)</td>
</tr>
<tr>
<td>Carry out dengue/vector borne surveillance in a village or college</td>
<td>14 (48.3)</td>
<td>13 (72.2)</td>
</tr>
</tbody>
</table>
Univariate analysis of public health skills with socio-demographic factors showed that students who are females, having graduation from Government College, being in 3rd year of PG, aged younger, mother's and father's professional education and occupation, have higher skill score than others, though the difference was not significant (Table 3).

**Discussion**

The medical education regulatory authority Medical Council of India (MCI) has defined essential competencies to be acquired at the end of PG education in Community Medicine. These include managerial and epidemiological skills in health care delivery system and national health programs, skills to identify community health needs, research methodology and abilities to impart training to medical and paramedical personnels (9). The present study was undertaken to assess some of these skills among the PG students of four Medical Colleges in Delhi.

Self assessment of communication and counselling skills by the PG students revealed that majority of them considered themselves to be competent to communicate with a family member of a patient and motivate eligible couple for contraceptive use. This finding is in contrast to the findings of study by Avan et al in 2005 in Pakistan (10). These high proportions give some credence to recognition by PG students of the importance of effective communication with the community and their patients. Further, interaction with people and their organization in

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**Table 3: Univariate Analysis of Total Skills Scores with Socio-demographic Variables**

<table>
<thead>
<tr>
<th>Variables</th>
<th>Score (mean±s.d.)</th>
<th>p value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gender</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Male (n=41)</td>
<td>15.71±4.11</td>
<td>0.185</td>
</tr>
<tr>
<td>Female (n=29)</td>
<td>17.00±3.79</td>
<td></td>
</tr>
<tr>
<td>Type of Medical College (from where MBBS done)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Government (n=34)</td>
<td>16.68±3.63</td>
<td>0.383</td>
</tr>
<tr>
<td>Private (n=36)</td>
<td>15.83±4.44</td>
<td></td>
</tr>
<tr>
<td>Year of Postgraduation</td>
<td></td>
<td></td>
</tr>
<tr>
<td>1st year (n=29)</td>
<td>15.34±4.05</td>
<td>0.203</td>
</tr>
<tr>
<td>2nd year (n=18)</td>
<td>16.28±3.01</td>
<td></td>
</tr>
<tr>
<td>3rd year (n=23)</td>
<td>17.35±4.49</td>
<td></td>
</tr>
<tr>
<td>Age</td>
<td></td>
<td></td>
</tr>
<tr>
<td>20-24 years (n=4)</td>
<td>17.00±2.16</td>
<td>0.206</td>
</tr>
<tr>
<td>25-29 years (n=61)</td>
<td>16.44±3.88</td>
<td></td>
</tr>
<tr>
<td>30-34 years (n=5)</td>
<td>13.20±5.85</td>
<td></td>
</tr>
<tr>
<td>Mother’s occupation</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Professional &amp; Semi-professional (n=9)</td>
<td>20.33±2.08</td>
<td>0.070</td>
</tr>
<tr>
<td>Others (n=61)</td>
<td>16.06±3.98</td>
<td></td>
</tr>
<tr>
<td>Father’s occupation</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Professional &amp; Semi-professional (n=44)</td>
<td>16.39±3.51</td>
<td>0.671</td>
</tr>
<tr>
<td>Others (n=26)</td>
<td>15.96±4.89</td>
<td></td>
</tr>
<tr>
<td>Mother’s education</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Professional (n=16)</td>
<td>16.44±3.46</td>
<td>0.829</td>
</tr>
<tr>
<td>Others (n=59)</td>
<td>16.19±4.18</td>
<td></td>
</tr>
<tr>
<td>Father’s education</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Professional (n=35)</td>
<td>16.94±3.24</td>
<td>0.145</td>
</tr>
<tr>
<td>Others (n=35)</td>
<td>15.54±4.58</td>
<td></td>
</tr>
</tbody>
</table>
urban and rural settings offers a wonderful opportunity for learning from people (9).

Epidemiology is an important part of Community Medicine and acquiring epidemiological competencies is essential for community physician (11). In the current study, only 58.6% considered themselves skilled enough to investigate an epidemic. Outbreak investigation is an opportunity to contain the epidemic and prevent future occurrence and is the best example for participatory and active learning of epidemiology. The finding highlights the need to focus on this skill among PG students during their training in epidemiology. Similarly planning evaluation of national health program or carrying out surveillance of a vector borne disease are essential components of epidemiology that offer opportunity to learn multiple things.

It is a ritual to establish public health laboratory to meet the requirements of MCI. However, at many places the laboratories have not been developed. Due to lack of infrastructure in the field areas, the skills in laboratory organization for organisms of public health importance is mostly lacking among PG students as was evident in the current study as well. Even though, the students were posted in public health laboratory of MCD and Microbiology Department of their respective colleges, many students found it hard to make smears for malaria diagnosis (41.4%) or test water sample for quality (72.9%). This finding points to the fact even though knowledge can be gained, learning is best experienced by doing, skills need to be practiced repeatedly to develop perfection (9).

In India, the requirement of occupational health manpower is accomplished by medical graduates, community medicine specialists, and public health specialists working as industrial medical officers (12). Thus, it is evident that PG student of Community Medicine should be trained in occupational health. However, in the present study, little less than half felt that they can do pre-placement examination of an industrial worker. This is probably is due to the fact occupational health training in India is most neglected in medical curriculum.

Limitations of Study

Since our study relied on self-reporting by the PG students, this may not correspond to their performance in actual clinical and community settings. The study also could not include reporting from community people and patients themselves, which might have strengthened the results and reflected their actual behavior. Also this was a cross-sectional study involving Medical Colleges in Delhi only. The variability in PG curriculum that exists in different medical colleges and universities of the country has not been investigated in the present study.

Nevertheless, the study highlights the skills which should be possessed by a PG student of Community Medicine. In the present study, though PG students assessed themselves to possess necessary skills on communication, counseling and health education, some skills related to occupational health, public health laboratory and epidemiology need to be inculcated among them with greater stress and indicate that these needed to be included in the PG Curriculum of the MD Program in Community Medicine. Further, there is a need to reform the curriculum in terms of competencies in several areas.

References


Neurodevelopmental Disorders: 
The Journey, the Dreams and their Realization

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All India Institute of Medical Sciences, New Delhi.

ABSTRACT

Neuro Developmental Disorders (NDDs) are associated with significant morbidity. This involves early identification of the disorder, the correct management of the disorder and associated disabilities. In India, the paucity of trained personnel and lack of knowledge about these disorders has been instrumental in inadequate management and recognition of these NDDs. The Child Neurology Division, Department of Pediatrics at All India Institute of Medical Sciences has made few noteworthy and meaningful contributions in these aspects: devising a DM curriculum for pediatric neurology, developing indigenous tools for diagnosing these NDDs and performing relevant research. These endeavors would go a long way in serving the children with NDDs.

Keywords: Neurodevelopmental disorders, autism, cerebral palsy, attention deficit hyperactivity disorder.

Introduction

Neuro Developmental Disorders (NDDs) constitute a significant proportion of morbidity handled by pediatric health care services. These include cerebral palsy, autism, attention deficit hyperactivity disorder (ADHD) and intellectual disability, epilepsy, autoimmune disorders including autoimmune encephalitis and neuromuscular disorders (including muscular dystrophy). A recent study estimated prevalence of ‘any NDD’ in 2–9 year old children in India to be 12.0% (95% CI: 10.9–13.2%), and 21.8% of these had more than one NDD (unpublished data). Thus, neurological disorders represent a significant cause of morbidity among children.

In addition, there has been a paucity of physicians trained in pediatric neurology and also, inadequate research focusing on disorders specific to our nation, which has further compromised care of these children. Hence, to overcome these lacunas, the Child Neurology Division, All India Institute of Medical Sciences has taken enormous steps over the past several years. These have been highlighted in subsequent sections.

DM Pediatric Neurology

In order to overcome the scarcity of physicians trained in Pediatric Neurology in our country, Pediatric Neurology program was
initiated in 2004, at All India Institute of Medical Sciences, under the Child Neurology Division as a part of Department of Pediatrics. Over the past 12 years, twenty-five residents have qualified as DM Pediatric Neurology. These residents are subjected to rigorous training under all aspects of clinical care as well as neuroradiology, neuropathology and other clinical and rehabilitative aspects of child neurology. The DM program at AIIMS has contributed to trained manpower not only in our country, but has also trained fellows from other SAARC nations, United States of America, France, and United Kingdom.

Research

There has been scarcity of research in pediatric neurological disorders from our country. Numerous national and international research projects have been undertaken at the Child Neurology Division, AIIMS, which contributed to the arena of epilepsy, autism, other NDD (like cerebral palsy), and autoimmune disorders. The division has been a pioneer in the country in developing dietary therapies for drug resistant epilepsy (1). The children whose epilepsy fails to be adequately controlled despite two antiepileptic drugs are defined to have drug resistant epilepsy (2). The therapeutic options for these children include epilepsy surgery or dietary therapy. The division has demonstrated the efficacy of these dietary therapies in drug resistant epilepsies using ketogenic diet, modified Atkins diet, and low glycemic index dietary therapy (3-7). In addition, it has contributed to the development of 2.5:1 ketogenic diet and proved its efficacy in institutional research (3). The division also, has been the frontrunner in diagnosing, managing and optimizing the treatment for autoimmune encephalitis (8), leukodystrophies (9), and acquired demyelinating disorders of CNS (10). Also, the research focusing on domiciliary management of intranasal and buccal midazolam has helped to standardize the management protocols for seizures at home (11). These researches have been instrumental in developing various therapeutic guidelines, and have also paved the way for various national programs including Rashtriya Bal Swasthya Karyakram (RBSK) (12).

Public Health Initiatives

Educating the parents and children and the society at large is an important aspect about management of any disease. The Child Neurology Division has played a pivotal role in educating the society about NDDs (Fig. 1),
autism (Fig. 2 to 4), and epilepsy by organizing regular public health lectures, television and radio shows/interviews, newspaper interviews and information materials. In addition, the division has published patient educational booklets on all the common pediatric neurological disorders. These are freely downloadable from http://www.aiims.edu/en/2014-12-24-07-16-28/neurology_education.html (Fig. 3) (13). Also the division has launched a helpline (with mobile number 9868399037 and email: autismhelp.pedsaiims@gmail.com, pedneuroaiims@yahoo.com) to facilitate help to parents of sick children.

Assisting Government Initiatives

The division was part of the INCLEN Collaborative study (with me being the Network Coordinator and Site PI), which estimated the prevalence of various neurological disorders across the country (14-17). This was instrumental in developing the convergence framework of RBSK (12). Also, the division has worked in close liaison with the Government of India for the certification of autism as a disability and issue of certificate for this disorder. The division in close collaboration with Ministry of Social Justice & Empowerment, Government of India, organized the workshop for training the master trainers for diagnosis of autism across the country in August-September 2016. This was aimed at formulating a uniform instrument for diagnosis of autism, and training health care professionals to use that tool and spreading awareness about the certification process.

Innovation

The division has been significantly contributed to several decisive innovations. These include use of skin biopsy to diagnose muscular dystrophy (especially collagen 6 related disorders) (18-20). Also, the division has revolutionized telephonic follow-up for disorders like neurocysticercosis, Lennox-Gastaut Syndrome, and Duchenne Muscular Dystrophy.

The division has also been pivotal in development of tools for diagnosis of various NDDs. Initially these tools were developed in collaboration with INCLEN (14-17). The tools
were later updated as per the latest guidelines. These AIIMS modified INDT tools for diagnosis of epilepsy, neuromuscular impairment, ADHD, and autism have been validated in institutional research. The tools for Autism Spectrum Disorder (https://play.google.com/store/apps/details?id=app.autism.sanyakhurana.diagnosticspectrumdisorder and https://itunes.apple.com/us/app/autism-spectrum-disorder-diagnostic/id1151524697?mt=8) and epilepsy (https://play.google.com/store/apps/details?id=com.weboutsourcing.childhoodepilepsy.childhoodepilepsydiagnostictool&hl=en and https://itunes.apple.com/in/app/childhood-epilepsy-diagnostic/id1078173463?mt=8) have been converted into a mobile application (Fig. 4) which is downloadable free of cost, while applications for other tools are under development.

**Microsite**

The Child Neurology Division of Department of Pediatrics at AIIMS launched a mobile-based website (m.nddworkshop2014.org) on 03-December-2014. This microsite was renamed as m.pedneuroaiims.org, in Dec 2015. This microsite deals in NDD including ADHD, Autism Spectrum Disorders, Intellectual Disability, Epilepsy, Learning Disability, Neuromuscular Disorders, Cerebral Palsy, Speech and Language Disorders, Hearing, and Vision Impairment. The microsite has various educational materials including videos, for doctors for clinical applications and for parents for home-based intervention (Fig. 5).

**Future**

The various initiatives mentioned above have laid the foundation for a healthier child neurology environment, where we can expect evidence-based management of neurological disorders among children. To build on these foundations, it is planned to perform high quality research covering all domains of neurology in collaboration with international leaders.

**Conclusion**

Neurological disorders are an enormous burden in terms of associated disabilities. In association most of these disorders are chronic and require patience and expertise for management. Various measures including
training physicians, research, innovations and public health initiatives will immensely contribute in improving the management of these children individually and also educate the society about the rights of these disabled children.

References


Biomedical Applications of Nanomaterials: Diagnosis and Therapy of Thrombotic Disorders

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ABSTRACT

We have employed unique properties of carbon-based as well as metallic nanomaterials to develop diagnostic / therapeutic devices targeted against thrombotic disorders. We have designed a novel graphene-based biosensor that can detect individuals with high coronary risk. Further, we describe an innovative strategy to ablate pathological thrombus in situ employing near-infrared laser-irradiated gold nanorods (photothermal therapy).

Keywords: Thrombus, biosensor, photothermal therapy, platelet-derived microparticles.

Introduction

Global status report of World Health Organization has categorized cardiovascular diseases, comprising of coronary artery disease, stroke, atherosclerosis and heart failure, as leading causes of death resulting in millions of casualties worldwide, followed by deaths due to diabetes. According to the WHO projection, the spectrum of cardiovascular diseases will be the largest cause of disability and death in India by 2020, thus replacing infectious diseases from that position. Although cardiovascular diseases affect people with advanced age, risk factors may appear relatively early. One such risk factor is circulating Platelet-derived Micro Particles (PMPs) in blood of patients suffering from myocardial infarction or peripheral arterial diseases (1). Elevated plasma levels of PMPs have been associated with enhanced risk of coronary heart disease in healthy individuals (2).

Platelet activation is central to the pathogenesis of arterial thrombosis. PMPs are membrane vesicles of less than 1 μm diameter released from stimulated platelets (3). They play significant role in haemostatic response and their presence in circulation represents serious procoagulant risk. Detection of PMPs at early stage of disease would aid in diagnosis, prevention and management of the pathology. Flow cytometry is the available technique for PMP detection but is associated with major drawbacks that include underestimation of PMP count, lack of universal standardization, time consuming experiments, high cost of the equipment and need for skilled operator. In contrast electrochemical biosensing stands far superior chance as an efficient and affordable tool for PMP detection.

'Biosensing' is an emerging concept based on amperometry, potentiometry and impedance.

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analysis, which detects transduction of biological events to electrochemical signals with high sensitivity (4). Nanoscale particles of gold, iron and silicon, as well as graphene and carbon nanotubes have been employed as immobilization matrices during fabrication of electrodes. Material immobilized on exposed surface of electrodes determines its specificity, which include enzymes, antibodies, proteins, aptamers or nucleotides depending upon the nature of the target. Although there has been attempt to detect microparticles using electrochemistry, the method does not differentiate between their cells of origin and types (5) and thus has limited medical application. Here we describe a simple, quick, sensitive and cost-effective method to detect PMPs circulating in blood of individuals with potential for point-of-care diagnostic at peripheral health care system. The novelty of this product lies in its ease of fabrication and detection method, as no electrochemical sensor has yet been reported which could facilitate quick screening and diagnosis of individuals at 'high-risk' for developing cardiovascular diseases, eliminating requirement of high end lab facilities and experienced technicians as needed in current PMP detection procedures.

Fig. 1. Schematic design of nano-biosensor for detection of PMPs depicting stepwise immobilization of GO and PAC1 antibody on electrode surface. Subsequent incubation of coated electrode with sample resulted in binding of platelet-derived microparticles, bearing active conformation of integrins α_{IIb}β_{IIIa}, to the sensor surface, which can be detected by impedance analysis.
We have designed a graphene oxide-based electrochemical biosensor for detection of PMPs, a major risk factor for arterial pro-thrombotic pathologies like acute myocardial infarction and ischemic stroke. Electrodes were fabricated with immobilized layers of graphene oxide and a specific antibody targeted against active conformation of integrin αβ3 on PMP surface. Results showed progressive rise in impedance in Nyquist plots with increasing number of PMPs in analyte. The sensor was highly specific for PMPs and did not identify microparticles originating from other cells. Blood obtained from patients diagnosed with acute myocardial infarction exhibited significantly higher values of impedance, consistent with larger number of circulating PMPs in these patients, as compared to samples from healthy individuals, thus validating biosensor as a specific, sensitive, label-free and cost-effective tool for rapid point-of-care detection of PMPs at bedside. Our biosensor is most ideal for mass population screening programs at periphery-level healthcare units with limited resources. It is aimed at early detection of individuals having higher imminent cardiovascular risk, as well as for routine analysis, which in turn would contribute to better management and survival of screened ‘high-risk’ subjects (6, 7).

Despite major scientific advances, non-communicable diseases like acute myocardial infarction, cerebral ischemic stroke, and deep vein thrombosis, underpinned with vascular blockage mediated by thrombus, remain major causes of death, and morbidity across the globe, incidence of which far exceeds that of cancer. Medication with fibrinolytic agents (tissue plasminogen activator, or tPA, streptokinase, and urokinase) are the widely used clinical practice that involves activation of plasminogen leading to proteolytic degradation of polymerized fibrin clot. However, fibrinolytic therapy is frequently associated with serious life-threatening complications like severe haemorrhage, embolism, haemorrhagic stroke, and reperfusion arrhythmias, and requires consistent physician supervision, and monitoring. Parenteral, or oral therapy would lead to generalized effect as it is not directed at localized clot, which, therefore, is another major limitation of this therapy. In this study, we have exploited biocompatibility, and photothermal attributes of NIR-active materials like gold nanorods (8) to dissolve fibrin clots at site of lesion, and restore lumen patency. The so-called photothermal therapy has been widely suggested as an anti-cancer measure to ablate solid tumours. This is the first report on application of photothermal therapy as an anti-thrombotic measure. The serine protease thrombin proteolytically degrades fibrinogen, the soluble blood protein component, to fibrin. Fibrin monomers are held together by non-covalent bonds constituting the nascent insoluble thrombus, which are further cross-linked by transglutaminase action of clotting factor XIII. Subjecting thrombus to photothermal heating would lyse non-covalent interactions, resulting in loosening, and downsizing of clot. Force of arterial, or venous fluid shear would disperse the remnants of loosened clot leading to drop in local concentration of polymerized fibrin, and relief from occlusion. Clot rarefication would also facilitate permeation of thrombolytics inside the thrombus mass.

Fibrinolytic therapy for arterial or venous thrombotic disorders warrants systemic administration of thrombolytics like streptokinase, which is associated with serious bleeding complications. In this study, we have provided proof-of-concept of photothermal ablation of thrombus Thrombi were generated in vitro either from purified fibrinogen or from plasma, or in vivo in murine blood vessels. Gold nanorods were added on fibrin-rich clots in vitro or targeted towards thrombi in situ in mice, followed by irradiation with a 808 nm near-infrared laser source at power density of 1.05 W/cm². Local rise in temperature (up to 55–65°C) was detected with an infrared thermal camera that leads to nearly 15% lysis of clot. This is the first report on application of photothermal therapy as an anti-thrombotic
Remarkably, addition of streptokinase has a multimodal additive effect in accelerating the photothermal lysis of thrombi (up to 40%) even at a dose significantly lower (by 30 to 50 times) than therapeutic concentration, thus minimizing life-threatening side effects and adverse complications. This combinatorial approach has great potential in bringing about lysis of pathological clots that can effectively overcome the drawbacks of existing therapies (9, 10). Thus, the synergism between photothermal therapy and chemotherapy with reduced bleeding risk opens possibility of a smart and safe multimodal thrombolytic regimen for future.

Fig. 2: Targeted photothermal therapy, when synergized with chemotherapy/thrombolytics at sub-therapeutic doses, exhibits great potential for the lysis of pathological clots with effective minimization of the life-threatening side effects and off-target adverse complications associated with existing therapy. This introduces the possibility of smart and safe multimodal thrombolytic regimen in the future.

Acknowledgements

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References


Skindex-29 to Determine Quality of Life and Emotional Factors in Dermatological Conditions

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ABSTRACT

Though rarely fatal, skin diseases are known to be associated with increased psychiatric morbidity and considerable impairment of quality of life (QoL). Health Related Quality of Life (HRQoL) in skin diseases can be assessed by generic or skin specific instruments. One hundred sixty patients with a range of dermatological diagnoses were studied on a cross sectional observational paradigm. Skindex-29, a skin specific instrument, was used to assess QoL, while anxiety and depression were assessed by Anxiety Status Inventory (ASI) and Depression Status Inventory (DSI), respectively. Sixty two patients (39%) had impaired QoL out of which 37(60%) had severe impairment. 11 patients (7%) had anxiety and 22(14%) had depression in the mild to moderate range. Gender, anxiety, depression and effect on appearance seem to be not related to impaired QoL. Further large scale studies are needed to ascertain the factors impinging on the QoL of dermatologically ill patients.

Keywords: Quality of life, anxiety, depression, skindex-29, skin diseases.
Introduction

Skin gets frequently damaged because it is directly in the 'firing line'. The 3000 and odd known diseases of skin form the most common health problems afflicting people worldwide (1). Some of these conditions are among the top ten leading causes of non-fatal disease burden (2). Though rarely life threatening, the effects of cutaneous disorders on patients' lives can be profound as in addition to the distress brought about by symptoms and lesions, they affect interpersonal interactions as the lesions as well as the treatment thereof affect the appearance of individuals (3). Prevalence of psychiatric disorders in dermatological patients is higher than that of certain medical conditions like cancer, cardiac and neurological disorders (4) but remain unrecognized and underestimated (5, 6). The interrelationships between skin and psychiatric disorders can be quite complex: Social isolation and stigma associated with skin conditions may result in psychiatric illnesses; poor hygiene in chronic psychiatric disorders can cause skin problems; psychiatric disorders may present as dermatological problems, (Delusional parasitosis, body dysmorpophobia) certain drugs used in dermatology may have secondary psychiatric problems and vice versa; psychiatric disorders may aggravate skin conditions and also may lead to poor help seeking behavior (7-9). Psychiatric morbidity in skin diseases therefore may adversely affect quality of life (QoL) in many ways.

Health-related quality of life (HRQoL), defined as the patients subjective evaluation of the influences of their current health status on their ability to achieve and maintain a level of overall functioning that allow them to achieve valued life goals and that are reflected in their general wellbeing (10), is considered as an essential outcome measure in addition to clinical parameters in skin diseases (11, 12). Upto a third of persons with skin diseases are known to suffer from psychopathology mainly in the form of anxiety and depression (13-16) but due to shortfall of required skills dermatologist's assessment of mental status of patients is known to be sensitive at 30% only (5) which underscores the importance of collaborative management of patients by dermatologists as well as psychiatrists. However, it would be wrong to assume that QoL is totally linked to psychiatric morbidity (17).

Psychiatric morbidity and HRQoL are fairly well investigated in specific dermatological conditions like Psoriasis (18, 19), Vitiligo (20, 21), Acne (22, 23), Alopecia (24, 25), Eczema (26) and Hansen's disease (27). There are just a few studies which investigated psychiatric morbidity and HRQoL in dermatological patients in general (8, 28-30). Somehow this field did not appear to have attracted investigators from India; two recent reviews from India hardly had any references of Indian studies (4, 17).

Material and Method

The study was conducted at a hospital attached to a medical college in central India on a cross sectional observational design. One hundred and sixty patients were selected at random from in-patient and outpatient population of the dermatology department. All the patients who were above 18 years of age and those who were willing to participate in the study only were accepted. Patients suffering from major medical illnesses, psychoses and significant cognitive impairments were excluded from the study. All the patients were furnished with written information about the nature of the study. The study was approved by the research and ethics committee of the institution.

All the skin diagnoses were made by dermatologists. Every patient was subjected to standard mental status examination. The following instruments were used to measure anxiety, depression and QoL:
**Anxiety Status Inventory (ASI):** This is a structured interview version of Zung's self-rating anxiety scale (31). 20 affective and somatic symptoms associated with anxiety are rated on a 4 point Likert scale (none, mild, moderate and severe). Maximum score is 80. Obtained score is converted into an index and graded into mild, moderate and severe. The test was translated into vernacular Hindi by translation and back translation by different experts and administered to 50 bilinguals. The correlation between English and Hindi version was found to good \((r=0.76, p<0.001)\). The Hindi version was utilized for the study.

**Depression Status Inventory (DSI):** This is a structured interview version of the Zung Self Rating Depression Scale (32). There are 20 items in this scale that explore the affective, psychological and somatic symptoms associated with depression. Each question is scored on a scale of 1 to 4. Maximum possible score is 80. Procedure similar to anxiety inventory was adapted to obtain Hindi version and correlation was found to be good \((r=0.79, p < 0.01)\). Obtained score in a given individual is converted into an index and graded into mild, moderate and severe grades.

**Skindex-29:** This is a skin specific QoL instrument devised by Chren (33). It has 30 items distributed into domains (symptoms, function and emotion). Each of the items is scaled on a 5 point Likert scale. All the responses are transferred to a linear scale (Never=0, rarely=25, sometimes=50, often=75 and all the time =100). Domain and overall scores can be computed. Overall score is the mean of the responses. Item number 18 is excluded from scoring as per the guidelines of the originators of the test. This test was also subjected to translation and back translation. Hindi version was found to be well correlated to the original English version \((r=0.86, p<0.001)\). QoL scores were graded into mild, moderate and severe grades as per Prinsen et al (34).

**Statistical Analysis:** Depression and anxiety scores were correlated with overall Skindex-29 scores by means of Pearson's product moment correlation test. Mean anxiety and depression scores of each grade of Skindex-29 were compared by means of one-way ANOVA followed by post hoc Tukey's test. One-way ANOVA was also used to compare the anxiety, depression and QoL scores in relation to symptom and diagnostic categories. Student's t-test was used to compare the anxiety, depression and Skindex-29 scores between genders. Domain scores were not subjected to analysis in this study.

**Results**

A total of 62 patients (39%) had impaired QoL out of which 15 had mild, 10 had moderate and 37 had severe grade impairments. Ninety eight patients reported no impairment in their QoL. Twenty two patients (14%) had depression while 11 (7%) had anxiety. Mean overall Skindex score was 25.27±20.21 (Mean±SD) while that of anxiety and depression was 32.3±8.43 and 39± 8.4, respectively. There was no significant difference \((t 0.39, p<0.05)\) between Skindex scores of males \((26.1±20.38)\) and females \((24.51±20.22)\). No significant difference was also found in the anxiety \((t 0.37, p<0.05)\), depression \((t 0.12, p<0.05)\) between genders. Overall Skindex-29 scores were found to be highly correlated to ASI \((r=0.29, p<0.001)\) and DSI scores \((r=0.37, p<0.001)\). We categorized patients into seven groups based on the predominant symptoms that they had, and compared their anxiety, depression and Skindex-29 scores. No significant differences in anxiety \((F 1.09, p>0.05)\) depression \((F 0.71, p>0.05)\) and Skindex scores \(( F 1.22, p>0.05)\) were found between patients with different symptoms (Table 1). ANOVA revealed uniform distribution of anxiety \((F 1.69, p>0.05)\) and depression \((F 1.02, p>0.05)\) scores across diagnostic categories. However, Skindex-29 scores were found to be differentially distributed \(( F 2.69, p<0.007)\) in various diagnoses: scabies patients had the lowest score (mean 13.8, SD
### Table 1: Anxiety, Depression, Skindex-29 QoL scores in-relation to Symptoms

<table>
<thead>
<tr>
<th>Symptoms</th>
<th>Anxiety</th>
<th>Depression</th>
<th>Skindex-29</th>
</tr>
</thead>
<tbody>
<tr>
<td>Eruptions (n=45)</td>
<td>26.6 (7.4)</td>
<td>31.6 (6.2)</td>
<td>24.9 (18.5)</td>
</tr>
<tr>
<td>Erythema (n=37)</td>
<td>27.1 (9.1)</td>
<td>33.4 (9.5)</td>
<td>27 (17.1)</td>
</tr>
<tr>
<td>Hyperpigmentation (n=32)</td>
<td>26.7 (8.2)</td>
<td>31.8 (6.5)</td>
<td>24.3 (21.1)</td>
</tr>
<tr>
<td>Hypopigmentation (n=20)</td>
<td>25.3 (7.2)</td>
<td>30.6 (5.9)</td>
<td>22.1 (18)</td>
</tr>
<tr>
<td>Itching (n=13)</td>
<td>21.8 (2.1)</td>
<td>29.3 (1.8)</td>
<td>22 (17)</td>
</tr>
<tr>
<td>Impaired Sensation (n=5)</td>
<td>30.4 (9.4)</td>
<td>31.8 (3.3)</td>
<td>45.2 (22)</td>
</tr>
<tr>
<td>Alopecia (n=8)</td>
<td>23 (4.85)</td>
<td>30 (20)</td>
<td>29.8 (1.8)</td>
</tr>
</tbody>
</table>

ANOVA Anxiety: $F=1.09$, Df = 6,153, $P=0.366$ NS
ANOVA Depression: $F=0.71$, Df = 6,153, $P=0.642$ NS
ANOVA QOL: $F=1.22$, Df = 6,153, $P=0.298$

### Table 2: Anxiety, Depression, Skindex-29 QoL scores in-relation to Diagnosis

<table>
<thead>
<tr>
<th>Diagnosis</th>
<th>N</th>
<th>Anxiety</th>
<th>Depression</th>
<th>Skindex-29</th>
</tr>
</thead>
<tbody>
<tr>
<td>Acne</td>
<td>38 (23.7)</td>
<td>29.9 (8.2)</td>
<td>32.2 (7.1)</td>
<td>26 (20.2)</td>
</tr>
<tr>
<td>Eczema</td>
<td>29 (18.1)</td>
<td>25.8 (6.2)</td>
<td>32.3 (5.6)</td>
<td>33 (19.8)</td>
</tr>
<tr>
<td>Tinea</td>
<td>27 (16.8)</td>
<td>28.4 (7.3)</td>
<td>32.6 (3.5)</td>
<td>27.6 (20.1)</td>
</tr>
<tr>
<td>Hansen disease</td>
<td>4 (2.5)</td>
<td>31.2 (10.9)</td>
<td>32 (3.8)</td>
<td>45.8 (25)</td>
</tr>
<tr>
<td>Psoriasis</td>
<td>17 (10.6)</td>
<td>30.7 (13.9)</td>
<td>34.6 (13)</td>
<td>38 (21.1)</td>
</tr>
<tr>
<td>Scabies</td>
<td>20 (12.5)</td>
<td>23 (3.2)</td>
<td>30 (2.3)</td>
<td>13.8 (9.9)</td>
</tr>
<tr>
<td>Vitiligo</td>
<td>6 (3.75)</td>
<td>22.1 (3.4)</td>
<td>26.1 (4.3)</td>
<td>16.1 (4.2)</td>
</tr>
<tr>
<td>Alopecia</td>
<td>8 (5)</td>
<td>23 (4.8)</td>
<td>30 (2)</td>
<td>29.8 (1.8)</td>
</tr>
<tr>
<td>Urticaria</td>
<td>11 (6.8)</td>
<td>22.3 (3.4)</td>
<td>30.4 (4.1)</td>
<td>21.7 (16)</td>
</tr>
</tbody>
</table>

ANOVA Anxiety: $F=1.69$, Df = 8,151, $P=0.099$ NS
ANOVA Depression: $F=1.02$, Df = 8,151, $P=0.429$ NS
ANOVA QOL: $F=2.69$, Df = 8,151, $P=0.007$ (Sig.)
Table 3: Skindex vs ASI

<table>
<thead>
<tr>
<th></th>
<th>Normal (N=98)</th>
<th>Mild (N=15)</th>
<th>Moderate (N=10)</th>
<th>Severe (N=37)</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Mean</strong></td>
<td>28.1</td>
<td>31.5</td>
<td>32.8</td>
<td>38.4</td>
</tr>
<tr>
<td><strong>SD</strong></td>
<td>7.9</td>
<td>5.7</td>
<td>8.7</td>
<td>14.3</td>
</tr>
<tr>
<td>F:4.69</td>
<td>P&lt;0.05</td>
<td>Df: 3,156</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Tukeys Test</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Normal vs mild</td>
<td>t=1.11</td>
<td>P&gt;0.05 (NS)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Mild vs moderate</td>
<td>t=0.15</td>
<td>P&gt;0.05 (NS)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Moderate vs severe</td>
<td>t=1.53</td>
<td>P&gt;0.05 (NS)</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

9.8) while people with Hansen’s disease had the highest impairment (mean 45.8, SD 25.0) (Table 2). Anxiety scores of patients with various grades of impairment of QoL though found to be non-homogenous (F 4.69, p <0.05) but multiple comparisons by Tukey’s test revealed no significant differences (Table 3). Depression scores of patients with different grades of QoL impairments were also found to be not uniform (F 4.37, p <0.05) but post hoc analysis by Tukey’s test revealed no significant differences (Table 4).

**Discussion**

The interaction of dermatological conditions on the psychosocial aspects of existence can be quite complex with matters impinging on such issues as satisfaction with appearance, body image, identity, socialization and sexuality with potential for restricted social interactions, reduced opportunities for employment (35, 29) and the obvious threat of, and sometimes actual, diminished opportunities for interpersonal attraction, friendship and mate selection. Clinicians routinely interpret measurements of physical functioning clinically and by investigations. Interpretations of the findings, in most medical illnesses, is fairly simple as they are compared against norms but in cases of skin disease due to intervening emotional factors, patient's experience with illnesses becomes a crucial factor in determining effectiveness of treatment (11). Suffering could be unique in many ways in skin conditions.

We attempted to assess QoL in a large sample of dermatologically ill by means of Skindex-29 which is considered to be an ideal tool for such an investigation (36). We analyzed only the overall scores in the present paper. We also attempted to find if the grades of severity of HRQoL correspond in any way to the grades of anxiety and depression measured by standard instruments. We believe this is the first time that such an investigation was carried out in India.

In the present study, we found more than a third of patients had reported impaired QoL, majority (60%) of which to a severe degree. HRQoL is a subjective estimate. Wide variations in scores are therefore possible as disease perceptions may vary from individual to individual and culture to culture. Skindex-29 scores ranging from 20 to 78 have been reported in literature (18, 29, 37-39).
Table 4: Skindex vs DSI

<table>
<thead>
<tr>
<th></th>
<th>Normal (N=98)</th>
<th>Mild (N=15)</th>
<th>Moderate (N=10)</th>
<th>Severe (N=37)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mean</td>
<td>36.5</td>
<td>39.0</td>
<td>39.5</td>
<td>43.9</td>
</tr>
<tr>
<td>SD</td>
<td>5.7</td>
<td>5.8</td>
<td>12.3</td>
<td>12.2</td>
</tr>
</tbody>
</table>

F: 4.37     P<0.05     Df: 3,156
Tukeys Test
- Normal vs mild  t=1.27  P>0.05 (NS)
- Mild vs moderate t=0.33  P>0.05 (NS)
- Moderate vs severe t=1.62  P>0.05 (NS)

The widely dispersed scores indicate differential effects but were not related to symptoms which we find somewhat counterintuitive as we imagined symptoms like pain, lack of pain, hyperpigmentation and hypopigmentation eruptions and so on will have different implications for the patients. Patients with different symptoms appear to be uniformly distressed (Table 1). The diagnostic label of the disease and chronicity seem to be significant determining factors as we found conditions like Hansens and psoriasis are associated with poor QoL in contrast to scabies, findings are in agreement with Kosaraju et al (40), Sanclemente et al (41). Effect of the disease lesions on appearance do not seem to be affecting the QoL much as we find obviously visible conditions like acne, vitiligo, tinea and urticaria are associated with fairly good QoL (Table 2). This finding, once again, is contrary to what is expected; perhaps analysis of domain scores will throw some light on this phenomenon.

In the present study, the prevalence of anxiety was found to be much less than that reported by Seyhan et al (8) (13.4%), Kumar et al (19) (52%), Jindal et al (42) (11.28%) and Aslam et al (43) (20%), Abebe et al (44) (37.4%). It is to be noted that they investigated individual skin conditions. Our study sample included all major and minor conditions hence it is possible the prevalence figures are getting averaged. We found significant correlation between anxiety scores and Skindex scores (r=0.29, p<0.001). Other investigators reported similar results (45-47). Though significant, anxiety is found to be contributing to only 8% of the variance of HRQoL. Further, we tried to assess if different grades of anxiety correspond to similar grades of impairment of HRQoL as reflected in Skindex grades. On the face of it ANOVA gives false impression of correspondence between anxiety and HRQoL scores. This is expected because the scores are correlated. But multiple comparison with Tukey’s test of the anxiety scores between normal and mild, mild and moderate and between moderate and severe grades of Skindex revealed no significant difference (Table 3).

The prevalence of depression (14%) in the present study also is less than that reported by Schmit and Ford (48) (31%), Basher et al (49) (34%), Sharma et al (50) (46.25%), Ponorovsky et al (30) (23.1%), Seyhan et al (8) (32%), Solgajova et al (51) (37.5%). Our finding of significant correlation (r 0.34, p<0.001) between depression scores and Skindex scores are similar to other studies (45-48). However, depression is seen to be
contribution to only 11% of the variance in Skindex-29 scores. ANOVA and posthoc Tukey’s test analysis of depression scores of different grades of Skindex scores yielded results similar to anxiety scores (Table 4). It appears depression and anxiety scores cannot serve as anchors to determine grades of severity of impairment of QoL.

Conclusion

The present study raises the familiar difficulty in interpreting HRQoL scores in patients with skin diseases (34, 52). The anchor based categories of Prinsen et al (34) may not serve the purpose in our settings. It also appears that skin diseases have different meanings to Indian patients than what is generally perceived. An anchor-based approach with appropriately worded anchor questions as well as a distribution based method Nijsten et al (53) with an additional normal sample may help in determining the grades of impairment of QoL. The cross sectional nature of the present study does not allow any inferences to be drawn about the responsivity of the scores to treatment. All categories of skin diseases are also not represented in adequate numbers in the present study. More importantly, what are the factors that have unique influence in determining QoL of dermatologically ill in our setting need to be explored fully to enable clinicians to chalk out appropriate strategies collaboratively to deliver comprehensive and holistic care to the clientele.

References


Restorative Therapies after Stroke: 
Drugs, Devices and Robotics

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ABSTRACT

Restorative therapies aim to improve outcome and function by promoting plasticity within a therapeutic time window between days to weeks to years. In this article, the mechanisms by which cell-based, pharmacological and robotic treatments stimulate endogenous brain remodelling after stroke, particularly neurogenesis, axonal plasticity and white-matter integrity are described with a brief outline of the potential of neuroimaging (fMRI) techniques. Stem cells aid stroke recovery via mechanisms depending on the type of cells used. Transplanted embryonic stem cells (ESCs), induced pluripotent stem cells (iPSCs), and neural stem cells (NSCs) can replace the missing brain cells in the Infarcted area, while adult stem cells, such as mesenchymal stem cells or multipotent stromal cells (MSCs) and MNCs, provide trophic support to enhance self-repair systems such as endogenous neurogenesis. Most preclinical studies of stem cell therapy for stroke have emphasized the need to enhance self-repair systems rather than to replace lost cells, regardless of the type of cells used. Noninvasive brain stimulation (NIBS) provides a valuable tool for interventional neurophysiology by modulating brain activity in a specific distributed, cortico-subcortical network. The two most commonly used techniques for noninvasive brain stimulation are transcranial magnetic stimulation (TMS) and transcranial direct current stimulation (tDCS). The article also discusses the potential role and current evidence for the use of pharmacological therapy, robotics and specific forms of physiotherapy regimes in optimizing stroke recovery. Neurorestoration is a concept that has been proven emphatically in several experimental models and clinical studies of stroke. Elucidating the underlying mechanisms of cell-based, pharmacological and rehabilitative therapies is of primary interest and crucial for translation of treatments to clinical use. The knowledge must provide an impetus for the development of superior, advanced and cost effective neuro restorative interventions that will enhance stroke recovery.

Keywords: Cerebral stroke, stroke therapy, functional neuroimaging.

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Introduction

Stroke is the second leading cause of mortality and morbidity with approximately half of them dependent on caregivers resulting in significant healthcare costs (1). Neuroprotection in the form of thrombolytic therapy is an inherent treatment for ischemia with only 5.2% receiving treatment in the window period (2). Acute stroke interventions for hemorrhagic stroke are limited. After much research focussed on acute neuroprotection, the National Institute of Neurological Disease and Stroke (NINDS) progress review group in 2006 and 2011 identified neurorestoration after stroke as a major priority for stroke research (3).

Restorative therapies aim to improve outcome and function by promoting plasticity within a therapeutic time window between days to weeks to years vis-a-vis reperfusion or neuroprotective drugs which act on salvageable brain (4). The aforementioned armamentarium includes growth factors, cell-based therapies, drugs and devices. Pharmacological treatment includes drugs that increase cGMP (e.g. phosphodiesterase 5 inhibitors, such as sildenafil and tadalafil), statins, erythropoietin, granulocyte-colony stimulating factor and minocycline (5).

There is an unmet need to design and improvise newer interventions to help patients repair and return to normal function. In this article, the mechanisms by which cell-based, pharmacological and robotic treatments stimulate endogenous brain remodelling after stroke, particularly neurogenesis, axonal plasticity and white-matter integrity are described with a brief outline of the potential of neuroimaging (fMRI) techniques (6).

Brain Insult: Repair and Recovery

The injury and recovery mechanisms after stroke have been extensively studied. The first epoch is related to acute injury and takes place in the first initial hours after stroke when changes in blood flow, edema, metabolism rate and diastasis occur. A second epoch is related to repair, which starts days after stroke and lasts for several weeks and is referred to as endogenous repair suggesting a golden period for initiating restorative therapies. A third epoch occurs weeks to months after stroke when spontaneous recovery gains have plateaued with residual deficits which may or may not be modifiable (7, 8).

Cell-based Interventions: Stem Cells

Stem cells can be defined as clonogenic cells that have the capacity to self-renew and differentiate into multiple cell lineages. They are divided according to the body's development process and their ability to form other cells (9). Totipotent stem cells are capable of giving rise to an entire organism and can be derived from fertilized oocytes and cells of the developing zygote up to the eighth cell stage. They have the potential to differentiate into derivatives of all germ layers (ectoderm, endoderm and mesoderm). Pluripotent stem cells can give rise to all tissue types, from any of the three embryonic germ layers, but unlike totipotent cells cannot give rise to an entire organism (10). These cells can give rise to different types of cells representing derivatives of two different germ layers, e.g. skin (ectoderm) and muscle (mesoderm). Multipotent stem cells are able to differentiate into multiple types of cells, but within one organ system (e.g. blood) (11). Progenitor cells differentiate into mature cells (e.g. endothelial progenitor cells) and can only divide a limited number of times, are laid between stem cells and fully differentiated cells (12-14).

The omnipresent nature of these cells, their clear role in neural tissue development, their presumed participation in repair and regeneration and the irrefutable success of bone marrow stem cell therapy have raised high expectations to cure diseases that have thus far proven resistant to conventional therapy such as...
stroke (15,16). The success of “bench to bedside” of cell transplantation in the last decade has seen a spurt of stem cell research in various pathological disorders, albeit all such clinical studies are/were phase 1/2 which aimed at safety and feasibility of cells.

**Human Umbilical Cord Blood Cells (UCB)**

These cells are derived from umbilical cord blood with a wondrous potential of differentiation into neural lineages (17). When exposed to nerve growth factor and retinoic acid, the derived umbilical cord blood cells produce progeny that shows positivity of neural and glial cells markers. A better understanding of these cells is needed before clinical transplantation studies ensue, although experimental data in animal models of stroke have shown functional benefits (18,19). However, biology of these cells is poorly understood, and it is likely that positive effects of these cells are related to their neurotrophic action, rather than actual neuronal circuitry formation.

**Immortalised Cell Lines**

These cell lines are derived by infecting neuroepithelial precursor cells from predefined CNS regions before terminal mitosis, with a retrovirus encoding an immortalizing oncogene view of the ethical difficulties in transplanting embryonic cells and technical problems in xenotransplantation, alternative sources of graft cells have been devised (20).

**Fetal Neural Stem Cells**

These cells maintain a normal karyotype for a significant number of passages in culture and can produce a large number of neurons and astrocytes and are harvested from the post-mortem human fetal brain. These possess a relatively high proliferative capacity without any evidence of tumorogenesis following transplantation (21).

**Adult Neural Stem Cells (NSC)**

Adult stem cells are multipotent stem cells found in developed organisms, which are used to replace cells that have died or lost function (22). NSCs are defined as undifferentiated cells that are able to self-renew as well as generate three major cell types of CNS: neurons, astrocytes and oligodendrocytes, signifying their pluripotent nature (23). They have been identified within many different organ systems, including bone marrow, brain, heart, skin and bone. Adult stem cells make up 1-2% of the total cell population within a particular tissue type. They are usually quiescent and held in an undifferentiated state until they receive a stimulus to differentiate (24).

**Bone Marrow Derived Cells**

These have hematopoietic and non-hematopoietic component, the former being abundant in bone marrow (25). Mobilized Peripheral Blood (MPB) is also a clinical source of heme cells, containing a mixture of hematopoietic stem and progenitor cells enriched with CD34 (26,27). These cells have the potential to regenerate the brain tissue by release of neurotrophic growth hormones. The other component of bone marrow contains mesenchymal stem cells or multipotent stromal cells (MSCs) described as colony-forming units (CFUs) that adhere to cell culture surfaces and can be differentiated into osteoblasts, adipocytes and chondrocytes (28-30). MSCs secrete interleukin-6 (IL-6), IL-7, IL-11, IL-12, leukemia inhibitory factor (LIF), macrophage-colony stimulating factor (M-CSF), stem cell factor (SCF) and flt-3 ligand (31).

**Induced Pluripotent Stem Cells (iPSCs)**

These cells are similar to human embryonic stem cells (ESCs) in morphology, proliferation, surface antigens, gene expression, epigenetic status of pluripotent cell-specific genes and telomerase activity (32). Adult human cells from skin were transformed to a pluripotent state using genetic engineering
Table: Clinical Trials of Stem Cells in Patient with Stroke

<table>
<thead>
<tr>
<th>Sr. No.</th>
<th>Study design control cell group</th>
<th>Characteristics of stroke</th>
<th>Manipulation (cell done)</th>
<th>Route</th>
<th>Efficacy</th>
<th>Adverse effects</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Autologous bone marrow mononuclear cells</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>1.</td>
<td>None : 5 patients 1-year f/u</td>
<td>Chronic Ischemic or ICH</td>
<td>Isolation using normal saline</td>
<td>IC</td>
<td>N/A</td>
<td>None</td>
</tr>
<tr>
<td>2.</td>
<td>None : 6 patients 6-month f/u</td>
<td>Subacute MCA infarct</td>
<td>Isolation using human albumin-containing normal saline (0.6-5x10^5)</td>
<td>IA</td>
<td>N/A</td>
<td>Seizure after 200 days</td>
</tr>
<tr>
<td>3.</td>
<td>None : 10 patients 6-month f/u</td>
<td>Acute Large MCA infarct</td>
<td>Isolation using human albumin-containing normal saline (0.6-5x10^5)</td>
<td>IV</td>
<td>Limited study design</td>
<td>None</td>
</tr>
<tr>
<td>4.</td>
<td>None : 20 patients 6-month f/u</td>
<td>Acute Nonlacunar infarct</td>
<td>Isolation using human albumin-containing normal saline (0.6-5x10^5)</td>
<td>IA</td>
<td>Limited study design</td>
<td>None</td>
</tr>
<tr>
<td>5.</td>
<td>40 : 60 patients 6-month f/u</td>
<td>Acute ICH</td>
<td>Isolation using normal saline (1.33x10^5)</td>
<td>IC</td>
<td>NIHSS and BI improved</td>
<td>None</td>
</tr>
<tr>
<td>6.</td>
<td>60 : 60 patients</td>
<td>Subacute MC/ACA infarct</td>
<td>Isolation using normal saline (2.8x10^5)</td>
<td>IV</td>
<td>B1 and mRS at day 180</td>
<td>Similar in the two groups</td>
</tr>
<tr>
<td><strong>Autologous bone marrow–derived mesenchymal stem cells</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>7.</td>
<td>25 : 5 patients 1-year f/u</td>
<td>Subacute Large MCA infarct</td>
<td>Ex vivo culture expansion using fetal bovine serum (1x10^5)</td>
<td>IV</td>
<td>B1 improved at 3 months</td>
<td>None</td>
</tr>
<tr>
<td>8.</td>
<td>36 : 16 patients 5-year f/u</td>
<td>Subacute Large MCA infarct</td>
<td>Ex vivo culture expansion using fetal bovine serum (1x10^5)</td>
<td>IV</td>
<td>mRS 0-3, increased in MSC group</td>
<td>None</td>
</tr>
<tr>
<td>9.</td>
<td>None : 12 patients 1-year f/u</td>
<td>Subacute to chronic Variable</td>
<td>Ex vivo culture expansion using autologous serum (1x10^5)</td>
<td>IV</td>
<td>Limited study design</td>
<td>None</td>
</tr>
<tr>
<td>10.</td>
<td>6:6 patients 24-week f/u</td>
<td>Chronic Ischemic or ICH</td>
<td>Ex vivo culture expansion using serum-free media (5-6x10^5)</td>
<td>IV</td>
<td>Modest increase in FM and mBI</td>
<td>None</td>
</tr>
<tr>
<td><strong>Allogeneic neural stem/progenitor cells</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>11.</td>
<td>None : 5 patients Terminated early</td>
<td>Chronic MCA infarct affecting striatum</td>
<td>Ex vivo culture expansion of NSCs obtained from primordial porcine striatum</td>
<td>IC</td>
<td>Limited study design</td>
<td>Seizure, aggravation of hemiplegia</td>
</tr>
<tr>
<td>12.</td>
<td>None : 8 patients 2-year f/u</td>
<td>Subacute to chronic MCA/ACA infarct</td>
<td>Ex vivo culture expansion of NSCs obtained from fetal brain</td>
<td>IC</td>
<td>Limited study design</td>
<td>Transient low-grade fever only</td>
</tr>
</tbody>
</table>

ACA: anterior cerebral artery, B1: Barthel index, FM: Meyer score, f/u: follow-up, IA: intra-arterial, IC: intra-cerebral, ICH: intra-cerebral hemorrhage, IV: intravenous, mBI: modified Barthel index, MCA: middle cerebral artery, mRS: modified Rankin Score, MSC: mesenchymal stem cell, N/A: not available, NIHSS: National Institutes of Health Stroke Scale, NSCs: neural stem/progenitor cells
techniques which could help generate patient and disease specific cells.

Stem cells aid stroke recovery via mechanisms depending on the type of cells used. Transplanted ESCs, iPSCs, and NSCs can replace the missing brain cells in the infarcted area, while adult stem cells, such as MSCs and MNCs, provide trophic support to enhance self-repair systems such as endogenous neurogenesis. Most preclinical studies of stem cell therapy for stroke have emphasized the need to enhance self-repair systems rather than to replace lost cells, regardless of the type of cells used.

**Clinical Trials of Stem Cell Research in Stroke**

Among all types of cells, bone marrow-derived stem cells are used frequently in clinical trials with stroke (Table 1). We have successfully transplanted bone marrow derived mononuclear and mesenchymal stem cells in chronic stroke (33-35).

The functional benefits after neural transplantation are likely to be mediated by one of the following mechanisms, i.e., neurotransmitters released from the graft tissue act on the afferent deprived limb of the post synaptic receptors, release of the neurotrophic / growth factors, Brain-derived Neurotrophic Factor (BDNF), Glial-derived Neurotrophic Factor (GDNF), Nerve Growth Factor (NGF) acting as local pumps to support cell function and to prevent cascade of apoptosis, regenerating neuronal population further prevents subsequent cell death, reestablishment of local interneuronal connections and synaptic connectivity between the host and graft, cell differentiation and integration, improvement of regional oxygen tension (36,37).

Our current stem cell trial investigates the paracrine mechanisms of mononuclear stem cells in chronic ischemic stroke, no serious adverse events were observed during the study.

There was no statistically significant clinical improvement between the groups (FM: 95% CI 15.2-5.35, p=0.25; mBI: 95% CI 14.3-4.5, p=0.31). Vascular Endothelial Growth Factor (VEGF) and BDNF expression was found to be greater in one group compared to other (VEGF: 442.1 vs. 400.3 pg/ml, p=0.67; BDNF: 21.3 vs. 19.5 ng/ml) without any statistically significant difference (38).

**Role of Noninvasive Brain Stimulation**

Noninvasive Brain Stimulation (NIBS) provides a valuable tool for interventional neurophysiology by modulating brain activity in a specific distributed, cortico-subcortical network (39, 40). Therapeutic utility of NIBS has been claimed in the literature for psychiatric disorders, such as depression, acutemania, bipolar disorders, schizophrenia, catatonia, post-traumatic stress disorder and neurologic diseases, such as stroke and parkinson's disease (41). The two most commonly used techniques for NIBS are Transcranial Magnetic Stimulation (TMS) and Transcranial Direct Current Stimulation (tDCS). TMS is a neurostimulation and neuromodulation application, whereas tDCS is a purely neuromodulatory intervention, both having their effect on the parameters of stimulation (42,43).

Hsu et al conducted a meta-analysis on eighteen studies involving 392 patients on efficacy of TMS post-stroke(44). A significant effect size of 0.55 was found for motor outcome (95% CI, 0.37-0.72) with subgroup analysis demonstrating more prominent effects on subcortical stroke (mean effect size~0.73; 95% CI, 0.44-1.02). Only 4 patients of the 18 articles included in this analysis reported adverse effects from rTMS (44). Low-frequency rTMS over the unaffected hemisphere may be more beneficial than high-frequency rTMS. Recent data suggest that intermittent theta-burst stimulation over the affected hemisphere might be a useful intervention (45).
Cicinelli et al (46,47) used focal TMS to map cortical representation via abductor digiti minimi (ADM) of the damaged hemisphere. It was observed that cortical excitability changed and there was an improvement in hand function. An ongoing research by our group which evaluates upregulation VEGF after acute ischemic stroke and its correlation with clinical recovery. It also examines the effects of rTMS (1Hz) and correlates the expression of VEGF in stroke patients. The results of this study are awaited.

Plautz (48) and Kleim et al (49) showed that cortical stimulation can reorganize movement representations to peri-infarct areas in primates and rats after ischemic lesions to their motor cortices. The importance of contralesional activation during motor tasks involving the recovering hand or arm is not clear. The effects seem to range from neutral or positive consequence such as adaptive neuroplastic process to negative maladaptation that may interfere with recovery.

In Transcranial Direct Current Stimulation (tDCS), low-amplitude direct currents are applied via scalp electrodes and penetrate the skull to enter the brain. Although the currents applied do not usually elicit action potentials, they modify the transmembrane neuronal potential and thus influence the level of excitability modulating the firing rate of individual neurons (50,51).

Lindenberg et al (52) conducted a study on twenty chronic stroke patients which were randomly assigned to receive 5 consecutive sessions of either a) bihemispheric tDCS (anodal tDCS to upregulate excitability of ipsilesional motor cortex and cathodal tDCS to down-regulate excitability of contralesional motor cortex) with physiotherapy or b) sham stimulation with physiotherapy. The improvement of motor function was significantly greater in the real stimulation group (20.7% in Fugl-Meyer and 19.1% in Wolf Motor Function Test scores) when compared to the sham group (3.2% in Fugl-Meyer and 6.0% in Wolf Motor Function Test scores) lasting for 1 week.

Anodal stimulation increases the spontaneous firing rate and the excitability of cortical neurons by depolarizing the membranes where as cathodal stimulation leads to neuronal hyperpolarization resulting in a decrease of the neuronal firing rate and excitability. It was found that anodal (facilitatory) tDCS stimulation of the ipsilesional hemisphere was associated with greater behavioral gains (better response time) as compared to cathodal (inhibitory) tDCS of the contralesional hemisphere and was associated with increased activation within ipsilesional M1 and PMd (53, 54). This pattern of activity was first shown in animals receiving stimulation via epidural or intracerebral electrodes. Therefore, facilitatory stimulation of the ipsilesional hemisphere may lead to gains in motor function by increasing activation within ipsilesional motor areas (55).

Our group, in a randomized placebo controlled trial is evaluating the efficacy of tDCS and fluoxetine on chronic stroke on balance and gait. The patients are randomized to four groups; real and sham tDCS and fluoxetine groups with dual task exercise regime administered to all patients. This is an ongoing study with results awaited.

Role of Mirror Therapy/ Virtual Reality

Mirror therapy was first described by Ramachandran in 1996 reporting its efficacy on pain reduction in arm amputees and later was claimed to alleviate hemiparesis after stroke. A pilot study confirmed the positive effects of mirror therapy on facilitation in upper limb hemiparesis after stroke (56). Mirror therapy is defined as an intervention that uses a mirror to create a reflection of the non-paretic upper or lower limb, thus giving the patient visual feedback of normal movement of the paretic limb. In a systematic review, fourteen studies were included with 9-121 participants. It was
found that mirror therapy had a significant effect on motor function (SMD 0.61; 95% CI: -0.22 to 1; p = 0.002; I² = 75%) (57). We also conducted a study with mirror therapy in chronic stroke patients using a web cam that captured the normal hand which was seen as affected in the laptop screen. Bilateral hand training were administered to patients and it was observed that MT improved hand function in FM and mBI scores along with an increased in laterality index (LI) in ipsilesional BA 4 and 6 (58).

**Role of Constraint Induced Movement Therapy (CIMT) and Electrical Stimulation**

CIMT has been investigated in 51 RCTs including 1784 patients with adult stroke, only 15 trials included patients within the first three months post-stroke. From systematic review (59), it is evident that original and modified versions of CIMT have a robust, clinically meaningful impact on patient's outcomes for arm-hand activities, self-reported hand use in daily life and basic ADL, making it one of the most effective interventions for the paretic limb post-stroke. It has also been reported that patients with poorer baseline behavior showed largest improvement on the wolf motor function test with increases in activation within the ipsilesional sensorimotor cortex during finger flexion/extension (60). Increased movement of the paretic upper extremity and decreased reliance on the non-paretic upper extremity are core features of CIMT (61).

Neuromuscular Electrical Stimulation (NMES) induces depolarization of peripheral neurons and subsequently elicits muscle contractions (62). It causes physiological changes after the stimulation, facilitating plastic changes during recovery and leading to improvement of voluntary functions. Functional Electrical Stimulation (FES) is one of the methods that uses electrical currents in stimulating the nerves connected to the paralyzed muscles in precise sequence and magnitude so that the outcome resembles functional tasks. FES aims to generate movements or functions, which mimic normal voluntary movements and therefore restore the functions (63). FES has some specific characteristics that makes it distinct from other forms of electrical stimulation. The frequency range of FES falls between 10 and 50 Hz and it directly stimulates the nerves or their motor points, not the muscle fibers. In 2002, Ada and Foongchomcheay (64) conducted a meta-analysis on the effect of electrical stimulation on shoulder outcomes after stroke. They showed that FES was superior to conventional therapy alone in the treatment of shoulder subluxation and arm motor function but was not effective in the treatment of pain early after stroke.

**Robotic Technology in Stroke**

The role of robotics in post-stroke rehabilitation has been investigated intensively. The robot-assisted rehabilitation of the upper limb in the acute and subacute post-stroke phase is successfully used as an alternative to conventional mobilization, resulting in effective conventional therapy. Masiero et al (65) hypothesized that an optimal robotic training protocol for acute and subacute stroke patients should be divided in two stages: initial additional robotic training (first stage) followed by substitution of part of the conventional therapy with the robotic exercise (second stage) (66). The introduction of robotic systems into clinical practice is useful in promoting a cost-effective use of human resources and the standardization of rehabilitation treatments. Hornby et al (67) performed a randomized controlled study comparing the effects of robot-assisted gait training that uses exoskeleton devices and manual facilitation that uses an assist on gait function in patients with chronic stroke. Another investigator also studied the usefulness of robot-assisted therapy in patients with subacute stroke in a multicenter randomized trial. They concluded that the diversity of conventional gait training interventions appeared to be more effective than robot-assisted gait training for improving walking ability (68). A study by Lo et al...
recruited 127 chronic stroke patients reported that robot-assisted therapy and conventional therapy produced similar amounts of improvement after 12 weeks of treatment and after 36 weeks of therapy, the robot-assisted therapy achieved greater motor improvement than did conventional therapy (69).

Pharmacological Agents

Pharmacological therapy post-stroke may seem merely a chimera, translation of drugs from the laboratory to the clinic should be performed with caution, failure of which from the bench to bed side transition will be devastating. For example erythropoietin (EPO) was demonstrated in multiple clinical studies to provide therapeutic benefit. Phase II clinical trial was unsuccessful and had to be terminated because of high mortality and adverse events.

Nitric Oxide (NO)

NO is an “endothelial-derived relaxing factor” which is involved in maintaining endothelial cell integrity, as well as participating in hemodynamic homeostasis (70). NO is also a potent activator of soluble guanylate cyclase, the enzyme that converts GTP to cGMP, the delivery of NO donor increases cGMP levels within both ischemic and non-ischemic rat brains, suggesting a permissive role for NO in neurogenesis. The increased expression of neuronal NO synthase within the Subventricular Zone (SVZ) during embryogenesis suggests an important role for the NO pathway in neurogenesis (71,72). In addition to enhancing cGMP levels by augmenting NO availability, cGMP levels may also be increased by inhibiting its metabolism by the Phosphodiesterase-5 (PDE5) enzyme (73). Animals treated with sildenafil, a PDE5 inhibitor post-stroke achieved significant and substantial increase in neurological functional recovery (74). It demonstrated improved cerebral blood flow (CBF), neurogenesis and synaptogenesis following experimental stroke, even when therapy was delayed for up to 1 week.

Gamma Amino Butyric Acid (GABA)

Recovery after stroke involves remapping of the neuronal circuitry in the regions adjacent to the site of injury or the peri infarct zone (75). A pharmacological approach to re-establish functional neuronal connections that are lost during stroke could enhance current physical rehabilitation therapies. It is also proven that inhibiting GABAergic signaling pathways after stroke can improve locomotor function, suggesting a therapeutic approach that is less time sensitive than acute reperfusion therapies.

Selective Serotonin Reuptake Inhibitors (SSRIs)

Animal studies suggest that SSRIs may be involved in neurogenesis and activation of cortical motor areas modulating neuronal plasticity (76). These drugs are essential in maintaining sleep rhythm, and neurotransmitters levels within the brain and have been tried in stroke rehabilitation trials. A single dose of citalopram can normalize the balance in cortical excitability, as measured by transmagnetic stimulation. Patients more than 6 months after stroke, in a single dose cross over experiment with citalopram, showed improvement in hand dexterity as measured by the nine-hole peg test (77). A single dose of fluoxetine given 2-3 weeks after stroke showed improved motor skills on the none-hole peg test, and increased activation of the affected side on functional resonance imaging. A meta-analysis of randomized controlled trials on stroke patients treated with SSRI compared to usual care or sham; identified 52 trials for analysis and it was found that these drugs are associated with an improvement in functionality, neurological impairment, disability and depression (78).

Minocycline

Minocycline is the second generation tetracycline derivative known to have anti-inflammatory effects independent of its antimicrobial action (79). Studies have shown
that minocycline prevents microglial activation, and has notable beneficial effects in animal models of global and transient focal cerebral ischemia. The proposed mechanisms of minocycline include anti-inflammatory effects, reduction of microglial activation, MMP reduction, NO production and inhibition of apoptotic cell death (80). In a randomized single blinded study, we studied the effects of oral minocycline (200 mg/day for 5 days) post-stroke versus placebo. Of 50 patients included in the trial, patients who received minocycline had better recovery in stroke outcome as noted on NIHSS, mBI and mRS scores (81).

Cerebrolysin

This is a peptide-based drug with potential to be used as a restorative agent. Multiple laboratories have demonstrated the safety and efficacy of this drug in the treatment of experimental stroke. It has been known to induce neurogenesis and angiogenesis in animal models of stroke and concomitantly enhances brain plasticity and recovery from stroke.

Niaspan

It is an extended release formulation of Niacin, proposed to be effective in reducing neurological deficits post-stroke by promoting axonal remodeling, angiogenesis and arteriogenesis (82). Niacin-induced increase in synaptic plasticity and axon growth may be mediated by the up-regulation in the BDNF–TrkB axis (Cui et al, 2010). In the mature nervous system, BDNF/TrkB plays an important role in regulating neuronal migration, differentiation, synaptic remodeling, and survival. Niacin treatment after stroke significantly increases BDNF/TrkB expression both in the ischemic brain and in primary cortical neuron (PCN) cultures (82). It has also been proven that combination of Niaspan with Simvastatin helped improve overall functional outcome significantly and decreased axonal damage and density (83).

Neuroimaging Provides Insight into Neuroplasticity

A wide number of neuroimaging methods exist for evaluating the state of Central Nervous System (CNS) function and structure after stroke. Brain function can be measured using functional Magnetic Resonance Imaging (fMRI), Positron Emission Tomography (PET), electroencephalography (EEG), magnetoencephalography (MEG), Transcranial Magnetic Stimulation (TMS) and near infrared spectroscopy (84). These techniques measure the volume of regional brain activation, the magnitude of activation and the balance of activation across hemispheres, often reported as a laterality index, during a task or at rest. Each technique has its merits and limits, MRI involves no isotopes and can also measure Cerebral Blood Flow (CBF) and angiography; PET can be used to measure CBF, metabolism, neurochemistry, and receptor kinetics; and TMS and MEG have temporal resolution at the millisecond level. Diffusion Tensor Imaging (DTI) is an MRI method for examining white matter integrity via measures such as Fractional Anisotropy (FA) enabling delineation of the anatomical connectivity of white-matter pathways (85).

An increasing number of studies have examined the mechanisms of spontaneous recovery after stroke. Studies have elucidated many of the cellular and molecular events, both near and remote from the lesion, that underlie spontaneous post-stroke improvements. These results are concordant with many of the findings from noninvasive neuroimaging methods in human subjects mentioned above.

Conclusion

Neurorestoration is a concept that has been proven emphatically in several experimental models and clinical studies of stroke. Elucidating the underlying mechanisms of cell-based, pharmacological and rehabilitative therapies is of primary interest and crucial for...
translation of treatments to clinical use. The knowledge must provide an impetus for the development of superior, advanced & cost-effective neurorestorative interventions that will enhance stroke recovery.

References


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