Expanding the comprehensive national neonatal screening programme in the United Arab Emirates from 1995 to 2011

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توسيع نطاق برنامج وطني شامل لتحري الولدان في الإمارات العربية المتحدة 1995-2011 هاجر الحوسني، محمد صلاح، هدى عثمان، حنان فرج، لبني الأسيوطي، دولا سعادة، جوزيف هرتكنت

ABSTRACT The national neonatal screening programme in the United Arab Emirates currently includes 16 disorders: congenital hypothyroidism, sickle-cell diseases, congenital adrenal hyperplasia, biotinidase deficiency and 12 amino acid, organic acid and fatty acid disorders. This paper reports data since the programme started in January 1995 up to December 2011 on the incidence of screened disorders and the molecular basis of positive screened cases. Screening used a combination of tandem mass spectrometry, molecular technologies and biochemical analysis. A total of 750 365 infants were screened and 717 babies saved from associated morbidity and/or mortality. The incidence of screened disorders were 1:1 873 for congenital hypothyroidism, 1:14 544 for phenylketonuria, 1:3 526 for amino acid, organic acid and fatty acid disorders, 1:9 030 for classical congenital adrenal hyperplasia, 1:8 300 for biotinidase deficiency, 1:2 384 for sickle-cell disease and 1:121 for sickle-cell traits. Coverage of neonatal screening in the population reached 95% in 2010.

Extension du vaste programme national de dépistage néonatal aux Émirats arabes unis de 1995 à 2011

RÉSUMÉ Le programme national de dépistage néonatal aux Émirats arabes unis couvre actuellement 16 maladies ou troubles : l'hyperthyroïdie congénitale, la drépanocytose, l'hyperplasie congénitale des surrénales, le déficit en biotinidase ainsi que 12 troubles des acides aminés, organiques et gras. L'article présente les données collectées, depuis le commencement du programme en janvier 1995 jusqu'en décembre 2011, sur l'incidence des troubles dépistés ainsi que la base moléculaire des cas positifs dépistés. La spectrométrie de masse en tandem, les technologies moléculaires et l'analyse biochimique ont été utilisées pour les besoins du dépistage. Au total, 750 365 nourrissons ont été dépistés et la morbidité et/ou mortalité associée a pu être évitée pour 717 bébés. L'incidence des maladies ayant fait l'objet d'un dépistage était la suivante : 1 : 1873 pour l'hyperthyroïdie congénitale, 1 : 14 544 pour la phénylcétonurie, 1 : 3526 pour les troubles des acides aminés, organiques et gras, 1 : 9030 pour la forme classique de l'hyperplasie congénitale des surrénales, 1 : 8300 pour le déficit en biotinidase, 1 : 2384 pour la drépanocytose et 1 : 121 pour les traits drépanocytaires. En 2010, la couverture de la population par le dépistage néonatal avait atteint 95 %.

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Introduction

The concept of screening newborns no longer refers only to the screening tests themselves but includes all of the elements essential for every neonate to have access to a screening system that is optimal in terms of quality and performance [1].

The United Arab Emirates (UAE) has a universal neonatal screening programme which started in January 1995 with screening for phenylketonuria (PKU) followed by the introduction of congenital hypothyroidism (CH) screening in January 1998. A sickle-cell disease (SCD) screening was begun as a pilot study in January 2002 before expansion in March 2005 to the national level. Congenital adrenal hyperplasia (CAH) screening also started as a pilot study in January 2005 before expansion to the national level in 2007. Screening for biotinidase deficiency was originally piloted in January 2010. Tandem mass spectrometry (MS/MS) technology expands the metabolic disorder screening panel and can reliably analyse approximately 50 metabolites in a single short-duration run of about 2 minutes and provide a comprehensive assessment from a single blood-spot specimen [2]. MS/MS was started in the UAE from March 2011 to screen for 12 amino acid, organic acid and fatty acid disorders.

The aim of the current study was to report data from the screening programme on the incidence of PKU, CH, CAH, SCD/trait, biotinidase deficiency, amino acid disorders, organic acid disorders and fatty acid disorders. It also aimed to identify the molecular basis of positive screened cases.

Methods

The sample forms were collected from the various maternal and child health centres throughout the UAE. The data from these forms and the results of the screening were statistically analysed in the computer section of the national screening programme. The screening protocol utilized a combination of MS/MS, molecular technologies and biochemical analysis.

Screening procedures

In the UAE, every baby born in hospital (around 80 000 babies with 99.9% hospital deliveries in 2010) is given a specific neonatal screening form and an information leaflet. The mother is asked to take her baby to the nearest screening centre, usually the maternal and child health (MCH) centre of that district. The baby attends the designated centre on the third day after birth (≥ 48 hours) and blood is collected by heel-prick onto filter paper (S&S 903) by technicians or trained nurses. For babies remaining in hospital for 3 days or more, blood is collected before departure. Quality assessment of laboratory results in the UAE is monitored by the schemes of the United Kingdom National External Quality Assessment Service and the United States Centers for Disease Control and Prevention. Any positive screening results are notified immediately to the central department of the MCH by the laboratory. The district MCH coordinator will be informed and will contact the parents.

For a baby screening positive for amino acid, organic acid and fatty acid oxidation disorders, CH, CAH and biotinidase deficiencies, further blood and urine samples are tested to confirm the diagnosis. For a baby screening positive for SCD or trait, blood is taken from the baby and family members before the age of 2 months for diseased and 4 months for trait cases to confirm the diagnosis and deliver genetic counselling. Testing of the parents for all haemoglobinopathies, including thalassaemia, may help establish the correct diagnosis in some infants.

Laboratory procedures

For screening for PKU (before March 2011), CAH, CH and biotinidase deficiency the Delfia* time-resolved fluorescence application (Wallac Oy) was used to determine levels of phenylalanine (PA), thyroid stimulating hormone (TSH), 17α-hydroxyprogesterone and biotinidase enzyme activity.

For detection of SCD the same filter papers were tested by high-performance liquid chromatography (Variant™, Bio-Rad).

Amino acid, organic acid and fatty acid oxidation disorders were detected by MS/MS (API 3200°, HVD/Perkin Elmer). At the time of the study 12 disorders were included in the screening: PKU, maple syrup disease, citrullinaemia type I, argininosuccinic acidaemia, isovaleric acidaemia, 3-methylcrotonyl-CoA carboxylase deficiency, hydroxymethylglutaric aciduria, beta-ketothiolase deficiency, glutaric acidaemia type I, propionic acidaemia, methylmalonic acidaemia and medium-chain acylcoenzyme A dehydrogenase deficiency.

Cut-offs

The following cut-offs were used for TSH levels (normal < 10 mU/L, borderline 10–25 mU/L, abnormal > 25 mU/L), PA levels (normal < 3 mg/dL, borderline 3–4 mg/dL, abnormal > 4 mg/dL), 17OHP levels (normal < 30 nmol/L, borderline 30–90 nmol/L, abnormal > 90 nmol/L) and biotinidase levels (normal > 30%, borderline 10%–30%, abnormal < 10%).

Molecular methods

Molecular diagnosis is done for specimens that show putative positive results following initial analysis to determine the presence or absence of specific mutations. This provides critical genetic data to confirm diagnosis and genetic counselling. The tests are done for all neonates, but for this study we selected the results of the UAE national babies to determine the UAE genetic profile and we excluded from our study the genetic

results of the other nationalities due to variable genetic profiles.

For PKU, the 13 exons and their exon-flanking intronic sequence of phenylalanine hydroxylase (PAH) gene were amplified by polymerase chain reaction (PCR) assay, and then singlestrand conformation polymorphism analysis for exons of the PAH gene and finally sequencing of gene exons was used to study uncharacterized PKU chromosomes [3]. The CAH confirmatory test was done by CAH strip assay covering 11 mutations in the CYP21A2 gene by reverse hybridization assay and a quantitative PCR approach to determine gene copy numbers [4]. For biotinidase deficiency, DNA sequence analysis was used to test for the presence of a mutation in all 4 exons of the biotinidase gene [5].

Results

Uptake of screening

Table 1 shows that the percentage uptake (coverage) of neonatal screening in the UAE increased from 50% in 1998 to reach 95% in 2010, with a rapid increase in 2003.

Congenital hypothyroidism

A total of 513 infants who screened positive for CH were further investigated and followed up; these included 129 infants with persistent borderline CH (TSH 10-25 mU/L) and 384 with abnormal results (TSH > 25 mU/L) (Table 2). Of the 129 borderline cases, 16 (12.4%) were confirmed CH. of whom 9 were defaulters. Of the infants with abnormal test results, 357 (92.9%) were confirmed CH. There were 60 transient cases that were normal on follow up. Among the transient cases, there were 14 cases with maternal history of autoimmune thyroid disease, 26 premature cases and 20 cases with unidentified causes. Of the 698 629 babies screened from the start (1 January 1988 to 31 December 2011) 373 were confirmed CH cases, an incidence of CH among the screened neonates of 1:1 873.

Phenylketonuria

Table 3 shows 132 neonates who screened positive for PKU; these were 75 with persistent borderline PKU (phenylalanine 3–4 mg/dL) and 57 abnormal babies (phenylalanine > 4 mg/dL)]. Of the 57 abnormal cases, 51 were confirmed as classic PKU (phenylalanine > 20 mg/dL, normal tyrosine and normal biopterin) and 1 case as PKU due to biopterin defect. Of the borderline cases, 61 (80.1%) were false positives. The incidence of PKU was 1:14 544.

Molecular DNA testing

A total of 7 different mutations (3 splicing mutations and 4 missense mutations) were detected by analysis of the PAH gene for 16 UAE national babies. The frequencies of the mutations were as follows: IVS2+5G>C (37.5%), R261Q (18.8%), IVS10-11G>A (12.5%), IVS9-2A>G (12.5%), R252Q (6.3%), P281L (6.3%) and L48S (6.3%).

Amino acid, organic acid and fatty acid disorders

Table 4 shows that 16 cases of amino acid, organic acid or fatty acid disorders

were detected by MS/MS from March 2011 until December 2011. There were 7 cases of hydroxymethylglutaric aciduria, 3 new cases of PKU (detected by MS/MS not by conventional methods) and 2 cases of glutaric acidaemia type I. The total incidence of these disorders was 1:353.

Sickle cell haemoglobinopathies

Table 5 shows that out of 5 498 positive screening results for sickle-cell haemoglobinopathies 231 neonates were SCD (FS), 80 had non-sickle haemoglobinopathy, 4481 were sickle haemoglobinopathy carriers (FAS) and 592 were haemoglobinopathy D carriers (FAD). The data showed that the incidence of SCD was 0.04%. It is noteworthy that 203 out of 231 SCD cases were confirmed as homozygous haemoglobin (Hb)S and only 28 cases were confirmed as sickle/beta-thalassaemia. The incidence of sickle-cell traits were 0.83% (1:121). For the non-sickle haemoglobinopathies, the incidence was 0.11% for HbD trait and 0.02% for HbC trait.

Congenital adrenal hyperplasia

Table 6 shows that out of 42 positive CAH screening results 41 babies were

Table 1 Coverage of the neonatal screening programme in the United Arab Emirates (1998 to 2010)

Year	No. of live births	% screened
1998	45 044	50
1999	49 075	61
2000	52 070	65
2001	53 485	67
2002	57 083	69
2003	60 249	89
2004	63 610	92
2005	66 192	94
2006	66 967	93
2007	67 789	94
2008	68 779	95
2009	76 366	95
2010	79 464	95

 ${\it Data \, source: \, Department \, of \, Preventive \, Medicine, \, Ministry \, of \, Health, \, United \, Arab \, Emirates.}$

Table 2 Follow up of newborns screening positive for congenital hypothyroidism (January 1998 to December 2011)

Variable	Borderline (TSH 10-25 mU/L)		Abnormal (TSH > 25 mU/L)		Total
	No.	%	No.	%	No.
Confirmed	16ª	12.4	357	92.9	373
Transient	49	38.0	11	2.9	60
False +ve	57	44.2	9	2.3	66
Failure to recall	7	5.4	7	1.9	14
Total +ve screening	129	100.0	384	100.0	513

^a9 cases were defaulters aged > 1 month when screened. TSH = thyroid stimulating hormone.

term deliveries and 1 was a preterm delivery. The data indicated that the number of false positive recalls for preterm deliveries (n = 175) was greater than for the term group (n = 77). The total incidence of classical CAH detected through newborn screening in the UAE was 1:9 030.

Molecular DNA testing

Of the 21 UAE national cases diagnosed as having CAH 18 were confirmed genetically as *CYP21* deficiencies, i.e. a mutation was found on both alleles: 10 were found to be homozygous for *I2* splice mutation; 6 had homozygous *CYP21A2* gene deletion; and 2 were heterozygous *I2* mutation. Three patients carried no *CYP21* mutations by the methods used.

Biotinidase deficiency

Three confirmed cases of biotinidase deficiency by enzymatic testing (1 profound and 2 partial cases) were discovered during the pilot study for

25 000 UAE national babies in 2010, an incidence of 1:8 300.

Molecular DNA testing

Molecular testing was done for only 2 confirmed cases. The first patient had partial biotinidase deficiency and 2 heterozygous point mutations were detected [C186Y (Cys186Tyr) and D444H (Asp444His)]. The second patient had profound biotinidase deficiency and 4 mutations were detected [2 homozygous mutations F403V (Phe403Val) and 2 homozygous mutations D444H (Asp444His)].

Discussion

Over the past 4 decades, many countries have developed screening programmes for newborns. An effective neonatal screening programme requires careful planning including education, administration, laboratory analysis, follow-up, management, evaluation and, most

importantly, the commitment of all involved [6].

Our data indicated that the percentage uptake of neonatal screening in the UAE increased from 50% in 1998 to reach 95% in 2010, with a rapid increase in 2003, although to levels still below the international coverage standard (99%) [7]. This improvement could be due to increased community awareness, a better recall system for defaulters, perinatal health education being combined with a breastfeeding programme and the recording of the screening status of the baby on the child welfare record along with vaccination information. The rapid increase in the rate of uptake in 2003 can be attributed to starting cooperation with Dubai medical district.

The incidence of CH among the screened neonates was 1:1 873. Of 373 confirmed cases, 357 had abnormal TSH level and 16 had persistent borderline TSH. Although high by international standards, the incidence of CH in the UAE was similar to that of other

Table 3 Follow up of newborns screening positive for phenylketonuria (PKU) (January 1995 to December 2011)

Variable	Borderline (PHA 3-4 mg/dL)		Abnormal (PHA > 4 mg/dL)		Total
	No.	%	No.	%	No.
PKU	-	0.0	52	91.2	52ª
Benign hyperplasia	11	15.1	5	8.8	16
Transient	3	4.8	-	0.0	3
False +ve	61	80.1	-	0.0	61
Total +ve screening	75	100.0	57	100.0	132

^{°51} cases were diagnosed as classical PKU and 1 case as PKU due to biopterin deficiency. PHA = phenylalanine.

Table 4 Amino acid, organic acid and fatty acid disorders in newborns screened by tandem mass spectrometry (March 2011 to December 2011) (n = 56 416)

Туре	No. of cases
Phenylketonuria	3
Maple syrup disease	1
Hydroxymethylglutaric aciduria	7
Glutaric acidaemia type I	2
Medium-chain acyl-coenzyme A dehydrogenase deficiency	1
Methylmalonic acidaemia	1
Isovaleric acidaemia	1
Total	16

countries in the region such as Oman (1:2 200) and Saudi Arabia (1:2 666) [8].

In our research, which used TSH as an indicator for the evaluation of iodine deficiency, the high incidence of CH was unlikely to be explained through iodine deficiency mechanisms [9]. Of the 513 cases that screened positive, 60 transient cases were normal on follow up. This was explained by maternal history and investigations of autoimmune thyroid diseases in 14 cases and the presence of prematurity in 26 cases. However, in 20 cases no specific cause was found. This deserves further investigation, e.g. obtaining an accurate clinical history of perinatal exposure to iodine, maternal antithyroid medication intake during pregnancy, TSH receptor antibodies and thyroid peroxidase

antibodies. These measures should help to decrease the recall rate when screening for CH and thus eliminate unnecessary stress to families, while contributing to a decline in the relatively high incidence of CH in the UAE [10].

There were 52 confirmed PKU cases over the 17-year period of screening (a relative incidence of 1:14 544); 51 were confirmed as classic PKU (phenylalanine > 20 mg/dL, normal tyrosine and normal biopterin) and 1 case as PKU due to biopterin defect. The incidence of PKU in the UAE was similar to that of other Gulf countries [11]. The high incidence of false positives within the borderline group might be due to genetic heterogeneity of hyperphenylalaninaemia and might indicate a need to adjust the cut-off point for the screening test [12].

A total of 7 different mutations with no novel ones were detected by analysis of the PAH gene for 16 UAE national cases. Three splicing mutations were identified, with the following frequencies [IVS2+5G>C (37.5%), IVS10–11G>A (12.5%) and IVS9-2A>G (12.5%)] and all were in the homozygous state. Four missense mutations were identified, with the following frequencies [R261Q (18.8%), R252Q (6.3%), P281L (6.3%) and L48S (6.3%)]. R261Q and R252Q were in the homozygous state but P281L and L48S were in the compound heterozygous state. The different mutations also contribute to a difference in phenotype and response to pharmacological doses of tetrahydrobiopterin [13].

There were 16 cases of amino acid, organic acid and fatty acid disorders detected by MS/MS over a 10-month period in 2011 (n = 56416 neonates screed), an incidence 1:3 526. The majority (7 cases) were hydroxymethylglutaric aciduria.

The figures for SCD (0.04%) and sickle-cell trait (0.83%) were lower than earlier studies done in the UAE, which recorded an incidence of 1.9% for sickle-cell trait [14]. A study in Saudi Arabia reported an incidence range of 2%–27% for sickle-cell trait and about 1.4% for SCD with the highest rates in the Eastern region

Table 5 Prevalence of confirmed sickle-cell diseases, non-sickle haemoglobinopathies and haemoglobinopathy carriers among newborns (January 2002 to December 2011) (n = 542 286)

Туре		Diseased			Carriers	
	No. identified	Incidence	Prevalence (%)	No. identified	Incidence	Prevalence (%)
Sickle Hb	231ª	1:238	0.04	4481	1:121	0.83
Non-sickle Hb	80	-	-	-	-	-
HbC	11	-	-	69	-	0.02
HbD	65	-	0.01	592	-	0.11
HbE	4	-	-	19	-	-
Unidentified band	-	-	-	11 ^b		-
Fetal Hb	-	-	-	15°	-	-

[°]Sickle-cell diseases: included 203 cases confirmed as homozygous SS and 28 cases confirmed as sickle/beta-thalassaemia; b3 diagnosed as HbO Arab by family study; cβ-thalassaemia and hereditary persistence of fetal haemoglobin.

Hb = haemoglobin.

Table 6 Follow up of newborns screening positive for congenital adrenal hyperplasia (CAH) from January 2007 to December 2011

Variable	Term	Preterm	Total
Screened population (no.)	367 436	11 836	379 272
Recalls (no.)	717	337	1 054
Recall rate (%)	0.2	2.9	0.28
CAH confirmed (no.)	41	1	42
False positive recalls (no.)	77	175	252

and the lowest in the Central region of the country [15]. Bahrain, on the other hand, had a rate of 11%–18% for sickle-cell trait, and an incidence of SCD of 2.1% among screened newborns [16]. Oman has reported an incidence of 10% for sickle-cell trait and about 0.4% for SCD [17].

The total incidence of classical CAH detected through newborn screening over a 10-year period in the UAE was 1:9 030. This figure is in agreement with older data from UAE derived from clinically detected cases [Azzam A, unpublished data, 2000] and with the high incidence of CAH reported in Saudi Arabia [18]. The high incidence of CAH in UAE and Saudi Arabia can be explained by the high rate of consanguineous marriages. The considerable false-positive recall rate for neonatal 17α-hydroxyprogesterone screening causes a substantial economical burden and emotional stress for parents and makes genetic testing an important confirmatory tool for early and reliable CAH diagnosis. More than 90% of classical CAH are caused by mutations in or deletions of the CYP21A2 gene encoding steroid 21-hydroxylase [19]. Eighteen of 21 local cases diagnosed as having CAH were confirmed genetically as CYP21 deficiencies with 10 being homozygous for I2 splice mutation and 6 homozygous CYP21A2 gene deletion. Two cases had heterozygous I2 mutation and, as the 2 cases were confirmed biochemically, we propose that these cases may be attributable

to compound heterozygosity of the *CYP21* gene with a rare or *de novo* mutation. Three patients carried no *CYP21* mutations by the method used and can also be attributable to other causes.

Three confirmed biotinidase deficiency cases by enzymatic testing (1 as profound and 2 as partial cases) were discovered during the pilot study for 25 000 UAE national babies in 2010, an incidence of 1:8 300. The disease incidence varies between countries, with higher incidences in countries with a high degree of consanguinity, such as Turkey, Saudi Arabia and Qatar (e.g. 1 in 12 607 screened babies in Qatar) [20]. The estimated international incidence of combined partial and profound biotinidase deficiency is 60 089 newborns (1:49 500 to 1:73 100) [21]. The first patient had partial biotinidase deficiency as confirmed by enzymatic and molecular testing. The molecular testing revealed 2 heterozygous mutations: C186Y mutations associated with profound biotinidase deficiency and D444H mutations associated with partial biotinidase deficiency. The presence of these heterozygous mutations in the biotinidase gene were consistent with a diagnosis of partial biotinidase deficiency. The second patient had profound biotinidase deficiency as confirmed by enzymatic and molecular testing. The molecular testing revealed 4 mutations: 2 homozygous F403V pathogenic mutations and 2 homozygous D444H mutations consistent with partial biotinidase deficiency. The presence of 2 double homozygous

mutations in the biotinidase gene was consistent with a diagnosis of profound biotinidase deficiency. No previous reports of similar cases were found in literature for the patient having 2 homozygous alterations (*F403V*; *D444H*). These alterations could be considered as novel mutations and could be related to a deletion of the exon, parental consanguinity, uniparental disomy or by chance alone. One way to address this issue is to test the parents and determine if they are carriers of the above mentioned alterations [22].

Conclusions

Based on the statistical data of the incidence of screened diseases in the UAE and the success achieved in early detection, treatment and follow-up, we emphasize the importance of expanding the programme capacity.

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