

One person can make a difference: identification of people with a rare genetic lung disease

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RSPH9 and *DNAH11*. Three of these variants were reported in a different cohort of Palestinians [8, 12–15]. Three novel candidate genes were also identified. To put this feat of work in perspective, a large PCD referral centre in the USA screens about 40–100 individuals for PCD per year.

Due to the high frequency of some alleles within specific families, likely due to founder mutations, the authors were able to use allele-specific PCR against mutations in *CCDC39* and *DNAAF11* for diagnosis. This option may allow a cheaper and more accessible alternative to commercial panels.

It is also noteworthy that despite the relatively high number of patients that were identified in this cohort, it is likely that many individuals with PCD may not have been identified. Genetic and TEM testing was highly selective and did not include patients with PCD like symptoms whose nNO measurements were >77 nL·min⁻¹ (the cut-off established by the ATS and ERS guidelines). Though this cut off is a good starting point, it is well established that many patients with PCD due to PCD genes (*e.g. RSPH1, GAS8, RPGR, CCNO, CCDC103, CFAP221, DNAH9, FOXJ1, GAS2L2, LRRC56, NEK10, SPEF2, STK36, HYDIN* or *TTC12*) have nNO values >77 nL·min⁻¹. This information suggests that the burden of disease may be even higher within the Palestinian population.

To improve the health of the people within the region, a consorted effort to establish centres of excellence in resource-restricted areas such as Palestine will be required. Such a response can improve the diagnosis and treatment of PCD, and other rare conditions such as genetic immunodeficiency diseases that have features overlapping PCD. The collaboration between the authors of this paper is one example of how this effort can be supported, starting by training talented individuals and providing the diagnostic support (locally or at a centralised centre overseas).

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References

- 1 Olbrich H, Haffner K, Kispert A, *et al.* Mutations in *DNAH5* cause primary ciliary dyskinesia and randomization of left-right asymmetry. *Nat Genet* 2002; 30: 143–144.
- 2 Omran H, Haffner K, Volkel A, *et al.* Homozygosity mapping of a gene locus for primary ciliary dyskinesia on chromosome 5p and identification of the heavy dynein chain *DNAH5* as a candidate gene. *Am J Respir Cell Mol Biol* 2000; 23: 696–702.
- 3 Shapiro AJ, Davis SD, Polineni D, *et al.* Diagnosis of primary ciliary dyskinesia. An official American Thoracic Society clinical practice guideline. *Am J Respir Crit Care Med* 2018; 197: e24–e39.
- 4 Rumman N, Fassad MR, Driessens C, *et al.* The Palestinian primary ciliary dyskinesia population: first results of the diagnostic and genetic spectrum. *ERJ Open Res* 2023; 9: 00714-2022.
- 5 Zlotogora J. Autosomal recessive diseases among Palestinian Arabs. J Med Genet 1997; 34: 765–766.
- 6 Khalaf-Nazzal R, Dweikat I, Maree M, et al. Prevalent MLC1 mutation causing autosomal recessive megalencephalic leukoencephalopathy in consanguineous Palestinian families. Brain Dev 2022; 44: 454–461.
- 7 Sirdah MM. Consanguinity profile in the Gaza Strip of Palestine: large-scale community-based study. Eur J Med Genet 2014; 57: 90–94.
- 8 Reish O, Slatkin M, Chapman-Shimshoni D, *et al.* Founder mutation(s) in the *RSPH9* gene leading to primary ciliary dyskinesia in two inbred Bedouin families. *Ann Hum Genet* 2010; 74: 117–125.
- 9 Alsaadi MM, Gaunt TR, Boustred CR, et al. From a single whole exome read to notions of clinical screening: primary ciliary dyskinesia and RSPH9 p.Lys268del in the Arabian Peninsula. Ann Hum Genet 2012: 76: 211–220.
- 10 Hammoudeh S, Gadelhak W, Janahi IA. Primary ciliary dyskinesia among Arabs: where do we go from here? *Paediatr Respir Rev* 2019; 29: 19–22.
- 11 Fassad MR, Shoman WI, Morsy H, *et al.* Clinical and genetic spectrum in 33 Egyptian families with suspected primary ciliary dyskinesia. *Clin Genet* 2020; 97: 509–515.
- 12 Monies D, Abouelhoda M, Assoum M, *et al.* Lessons learned from large-scale, first-tier clinical exome sequencing in a highly consanguineous population. *Am J Hum Genet* 2019; 105: 879.
- 13 Merveille AC, Davis EE, Becker-Heck A, et al. CCDC39 is required for assembly of inner dynein arms and the dynein regulatory complex and for normal ciliary motility in humans and dogs. Nat Genet 2011; 43: 72–78.
- 14 Horani A, Ferkol TW, Shoseyov D, *et al. LRRC6* mutation causes primary ciliary dyskinesia with dynein arm defects. *PLoS One* 2013; 8: e59436.
- **15** Castleman VH, Romio L, Chodhari R, *et al.* Mutations in radial spoke head protein genes RSPH9 and RSPH4A cause primary ciliary dyskinesia with central-microtubular-pair abnormalities. *Am J Hum Genet* 2009; 84: 197–209.