The University of Chicago Genetic Services Laboratories



5841 S. Maryland Ave., Rm. G701, MC 0077, Chicago, Illinois 60637

Toll Free: (888) UC GENES (888) 824 3637 Local: (773) 834 0555 FAX: (773) 702 9130

ucgslabs@genetics.uchicago.edu

dnatesting.uchicago.edu

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Prader-Willi Syndrome Testing

Clinical Features:

Prader-Willi syndrome (PWS) [OMIM #176270] is a genetic disorder which causes hypotonia and poor feeding in infancy, followed by the development of hyperphagia and subsequent obesity. Physical characteristics of PWS also include short stature, small hands and feet, and a characteristic facial appearance consisting of a thin upper-lip, down-turned mouth, dental crowding, and almond shaped eyes. Developmental milestones are delayed, and learning disabilities are always present, but may vary in severity. Behavioral problems include temper tantrums, obsessive compulsive tendencies, and skin-picking. Psychosis occurs in 5-10% of patients as young adults (1). Individuals with PWS do not undergo spontaneous pubertal development and are infertile (1, 2).

Inheritance:

PWS is caused by the absence of paternally expressed, maternally silenced genes in the imprinted region located at 15q11-q13. This can occur by one of several mechanisms including a *de novo* paternal deletion in this region, maternal uniparental disomy (UPD), an imprinting center defect, or a paternal chromosomal translocation. Mutations in the paternal copy of the *MAGEL2* gene located at 15q11.2 have also been described in patients with classic PWS and patients with PWS-like features (3). The recurrence risk depends on the mechanism involved and may be up to 50%. PWS affects approximately 1 in 25,000 births and displays no ethnic or gender preference (2, 4).

Molecular Genetics:

- Approximately 70% of individuals with PWS have a de novo deletion of 15q11-q13 on the paternally contributed chromosome, corresponding to a less than 1% recurrence risk (2).
- Approximately 25% of PWS is due to maternal UPD15, corresponding to a less than 1% recurrence risk (2).
- ≥ 2-5% of patients have an imprinting center (IC) abnormality of which 10-40% are deletions of the IC region. A recurrence risk of up to 50% applies to the IC deletion group and a low recurrence risk of less than 1% applies to the remainder of the IC abnormality group (2).
- Less than 1% of PWS is due to a paternal chromosome 15 translocation, which may result in a recurrence risk of up to 25% (2).
- De novo truncating mutations in the paternally inherited copy of MAGEL2 have been associated with classic PWS or PWS-like features (3). The proportion of PWS associated with MAGEL2 mutations is unknown, but is predicted to be low.

Additional Resources:

Prader-Willi Syndrome Association

5700 Midnight Pass Road, Suite 6 Sarasota, FL 34242 Phone: 800-926-4797 email: national@pwsausa.org www.pwsausa.org/index.html

Test methods:

We recommend methylation-specific (MS)-MLPA as the initial test for PWS. This testing will identify patients with abnormal methylation, large deletions and imprinting center deletions. Those patients with abnormal methylation, but no deletion, should pursue UPD testing for UPD15. For patients with normal MS-MLPA results, sequencing of the *MAGEL2* gene can be considered.

Methylation-specific MLPA (MS-MLPA)

Sample specifications: 3-10 cc of blood in a purple top (EDTA) tube

Cost: \$525 CPT codes: 81331 Turn-around time: 4 weeks

Microsatellite analysis for UPD15 testing

Sample specifications: 3-10 cc blood from patient

and BOTH parents in purple top EDTA tubes

Cost: \$540 (total for a patient's and both parents' blood samples)

CPT codes: 81402 Turnaround time: 2 – 4 weeks

Imprinting center deletion analysis

Sample specifications: 3-10 cc of blood in a purple top (EDTA) tube

Cost: \$450 CPT codes: 81403 Turn-around time: 4 weeks

MAGEL2 sequence analysis

Sample specifications: 3-10 cc of blood in a purple top (EDTA) tube

Cost: \$1,575 CPT codes: 81403 Turn-around time: 4 weeks

Results:

Results, along with an interpretive report, will be faxed to the referring physician. Additional reports will be provided as requested. All abnormal results will be reported by telephone.

References:

- 1. Chen C, Visootsak J, Dills S et al. Prader-Willi syndrome: an update and review for the primary pediatrician. Clin Pediatr (Phila) 2007: 46: 580-591.
- 2. Driscoll D, Miller J, Schwartz S et al. Prader-Willi Syndrome. In: Pagon R, Bird T, Dolan C, eds. GeneReviews [Internet]. Seattle: University of Washington, 1998.
- 3. Schaaf CP, Gonzalez-Garay ML, Xia F et al. Truncating mutations of MAGEL2 cause Prader-Willi phenotypes and autism. Nat Genet 2013: 45: 1405-1408.
- 4. Whittington JE, Holland AJ, Webb T et al. Population prevalence and estimated birth incidence and mortality rate for people with Prader-Willi syndrome in one UK Health Region. J Med Genet 2001: 38: 792-798.

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