

Central congenital hypothyroidism caused by a novel mutation, C47W, in the cysteine knot region of TSH $\beta$

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Supplemental Table 1. List of known genes related to thyroid disorders.

Supplemental Figure 1. Protein modeling in silico was performed for TSH $\beta$ , wild type (left) and mutant (right). Mutation site (in red) causes significant change in the three-dimensional predicted structure as shown. Methodology for modeling: Kallberg M, Wang H, Wang S, Peng J, Wang Z, Lu H, Xu. Template-based protein structure modeling using the RaptorX web server. Nat Protoc 2012;7:1511-1522. Ma J, Peng J, Wang S, Xu J. A conditional neural fields model for protein threading. Bioinformatics 2012;28:i59-i66. Ma J, Wang S, Zhao F, Xu J. Protein threading using context-specific alignment potential. Bioinformatics 2013;29:i257-i265.

Genes Related to thyroid disorder								
<i>AADAT</i>	<i>DUOX1</i>	<i>GNAS</i>	<i>KMT2D</i>	<i>POU1F1</i>	<i>RYR2</i>	<i>SLC30A10</i>	<i>THRB</i>	<i>TTR</i>
<i>ALB</i>	<i>DUOXA1</i>	<i>HHEX</i>	<i>NCOR2</i>	<i>PROPI</i>	<i>SALL1</i>	<i>SLC5A5</i>	<i>TPO</i>	<i>TUBB1</i>
<i>ALMS1</i>	<i>DUOX2</i>	<i>HOXA3</i>	<i>NKX2-1</i>	<i>PSMA1</i>	<i>SECISBP2</i>	<i>SLCO1C1</i>	<i>TRH</i>	<i>UBR1</i>
<i>ATXN2</i>	<i>DUOXA2</i>	<i>IGSF1</i>	<i>NKX2-5</i>	<i>PSMA3</i>	<i>SERPINA7</i>	<i>STAMBP</i>	<i>TRHR</i>	<i>VAV3</i>
<i>CDC48</i>	<i>EXOSC2</i>	<i>IRS4</i>	<i>NKX2-6</i>	<i>PSMD2</i>	<i>SLC16A2</i>	<i>TBL1X</i>	<i>TRIP11</i>	
<i>DIO1</i>	<i>FGF8</i>	<i>IYD</i>	<i>NTN1</i>	<i>PSMD3</i>	<i>SLC17A4</i>	<i>TBX1</i>	<i>TRIP12</i>	
<i>DIO2</i>	<i>FOXE1</i>	<i>JAG1</i>	<i>P4HB</i>	<i>PTH1R</i>	<i>SLC26A4</i>	<i>TG</i>	<i>TSHB</i>	
<i>DIO3</i>	<i>GLIS3</i>	<i>KDM6A</i>	<i>PAX8</i>	<i>PTRH2</i>	<i>SLC26A7</i>	<i>THRA</i>	<i>TSHR</i>	

## TSHB Protein Modeling



Wild Type



c.T141G p.C47W