

Appendix 7

Northern blot results and transcript sizes (kb). The most dominant transcripts are in bold.

| <i>Gene</i> | <i>heart</i> | <i>brain</i> | <i>placenta</i> | <i>lung</i> | <i>liver</i> | <i>skeletal muscle</i> | <i>kidney</i> | <i>pancreas</i> |
|----------------|---------------------------------|--------------------------|-------------------------------|----------------------|--|-------------------------------|-----------------------------|---------------------------|
| AIF1_6p21.33 | - 3.0 1.25 0.6 | - 3.0 - | - 3.0 1.25 - | - 3.0 - | - - | 5.0 3.0 - | - 3.0 - | - 3.0 - |
| AIF1-L_9q34.12 | 3.4 | 3.4 | 3.4 | 0 | 0 | 0 | 3.4 | 0 |
| BRD2_6p21.32 | 4.6 3.8 | 4.6 3.8 | 4.6 3.8 | 4.6 3.8 | 4.6 3.8 | 4.6 3.8 | 4.6 3.8 | 4.6 3.8 |
| BRDT_1p22.1 | 7.0 - - | 7.0 - - | 7.0 - - | 0 | 7.0 - - | 7.0 - - | 7.0 - - | 7.0 4.0 3.5 |
| BRD3_9q34.2 | 6.5 | 6.5 | 6.5 | 6.5 | 6.5 | 6.5 | 6.5 | 6.5 |
| BRD4_19p13.12 | 6.0 4.4 | 6.0 4.4 | 6.0 4.4 | 6.0 4.4 | 6.0 4.4 | 6.0 4.4 | 6.0 4.4 | 6.0 4.4 |
| C4_6p21.33 | 0 | 0 | 0 | 0 | 5.4 | 0 | 5.4 | 0 |
| C5_9q33.2 | 0 | 0 | 0 | 0 | 6.0 5.0 4.2 1.6 1.0 | 0 | 5.0 - - - - | 5.0 4.2 - - - |
| C3_19p13.3 | 0 | 0 | 0 | 0 | 5.0 | 0 | 0 | 0 |
| CLIC1_6p21.33 | 1.25 1.1 | 0 | 1.25 1.1 | 1.25 1.1 | 1.25 1.1 | 1.25 1.1 | 1.25 1.1 | 1.25 1.1 |
| CLIC4_1p35.3 | 4.0 | 0 | 4.0 | 4.0 | 4.0 | 4.0 | 4.0 | 0 |
| CLIC3_9q34.3 | - 4.4 - - | 0 | 5.5 - 2.6 0.7 | - - - 0.7 | - - - 0.7 | 4.4 - - - | 4.4 - - - | 4.4 - - - |
| CLIC5_6p21.1 | 0 | 2.7 2.4 | 0 | 2.7 | 2.7 | 0 | 2.7 2.4 | 2.7 |
| CLIC6_21q22.12 | 6.0 3.8 3.0 2.3 | 0 | 0 | - 3.8 3.0 - | - - - 0 | 6.0 3.8 - 0 | - - - 0 | - - 3.0 - |
| CLIC2_Xq28 | 1.7 1.25 | 0 | 1.7 1.25 | 1.7 1.25 | 1.7 1.25 | 1.7 1.25 | 1.7 1.25 | 1.7 1.25 |
| GPX5_6p22.1 | - 0.8 0.6 | - 0.8 - | - 0.6 | 0 | - 0.8 0.6 | 3.8 0.8 0.6 | 0 | 0.8 0.6 |
| GPX4_19p13.3 | 4.4 2.6 2.0 0.9 | - - - 0.9 | - 2.6 2.0 0.9 | - - - 0.9 | 4.4 2.6 2.0 0.9 | 4.4 - 2.0 0.9 | - - 2.0 0.9 | - - - 0.9 |
| GPX1_3p21.31 | 0.8 | 0.8 | 0.8 | 0.8 | 0.8 | 0.8 | 0.8 | 0.8 |
| GPX3_5q33.1 | - 0.7 | 0 | - 0.7 | - 0.7 | 1.8 0.7 | 1.8 0.7 | - 0.7 | - 0.7 |
| GPX2_14q23.3 | 0 | 0 | 0 | 0 | 0 | 0 | 0.9 | 0 |
| NOTCH4_6p21.33 | 7.8 6.8 2.4 | - - - | 7.5 6.8 - | 7.5 6.8 - | - - 2.4 | - - 2.4 | - - - | - - 2.4 |

| | | | | | | | | |
|-----------------|------------------------------------|-------------------------------------|--------------------------------|------------------------------|-------------------------------------|-------------------------------|---------------------------------|-------------------------------|
| | - 0.8 | 1.6 - | - 0.8 | - - | - 0.8 | - 0.8 | - - | 1.6 0.8 |
| NOTCH2_1p11.2 | 10 1.9 | 10 - | 10 1.9 | 10 1.9 | 10 1.9 | 10 1.9 | 10 1.9 | 10 1.9 |
| NOTCH1_9q34.3 | 9.3 | 0 | 9.3 | 0 | 9.3 | 9.3 | 9.3 | 0 |
| NOTCH3_19p13.12 | 8 | 8 | 8 | 0 | 8 | 8 | 8 | 8 |
| PBX2_6p21.33 | 3.2 | 3.2 | 3.2 | 3.2 | 3.2 | 3.2 | 3.2 | 3.2 |
| PBX1_1q23.3 | 7.5 - 2.9 - | 7.5 - 2.9 - | 7.5 - 2.9 - | 7.5 - 2.9 - | 7.5 - 2.9 - | 7.5 4.4 2.9 - | 7.5 - 2.9 - | 7.5 4.4 2.9 - |
| PBX3_9q33.3 | 2.5 2.2 1.4 | 2.5 - - | 2.5 - - | 2.5 - - | 2.5 - - | 2.5 2.2 - | 0 | 2.5 2.2 - |
| PBX4_19p13.11 | - 4.5 1.4 1.2 | - 4.5 - - | - 4.5 1.4 1.2 | 0 | - 4.5 1.4 1.2 | 6.0 4.5 1.4 - | - 4.5 - - | - 4.5 1.4 - |
| RXRB_6p21.32 | 7.7 2.8 1.7 | - 2.8 - | - 2.8 - | 0 | - - 1.7 | 7.7 2.8 1.7 | - 2.8 - | - 2.8 - |
| RXRG_1q23.3 | 1.7 | 0 | 2.8 0.9 | 0 | 1.7 1.5 | 1.7 | 0 | 1.7 1.5 |
| RXRA_9q34.2 | 5.4 | 5.4 | 5.4 | 0 | 5.4 | 5.4 | 0 | 0 |
| TUBB_6p21.3 | - 4.8 4.0 2.5 1.7 - | - - - - 1.7 - | 5.1 - - - 1.7 - | - - - - 1.7 - | - - 4.8 4.0 1.7 1.35 | - - - - 1.7 - | - - - - 1.7 1.35 | - - - - - - |
| TUBB4QL_10p15.3 | 0 | 2.8 | 0 | 0 | 0 | 0 | 0 | 0 |
| TUBB4_16q24.3 | 1.7 1.3 | 1.7 1.3 | 1.7 1.3 | 1.7 1.3 | 1.7 1.3 | 1.7 1.3 | 1.7 1.3 | 1.7 1.3 |
| TUBBL_18p11.1 | 1.3 | 1.3 | 1.3 | 1.3 | 1.3 | 1.3 | 1.3 | 1.3 |
| TUBB5_19p13.3 | 0 | 2.3 | 0 | 0 | 0 | 0 | 0 | 2.3 |
| TUBB1_20q13.32 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |