**Additional file 5. Genes tested by qRT-PCR in retina and RPE\*.** Genes are reported with their symbols (in parenthesis alternative symbols), descriptions, main functions, and TaqMan assay numbers (ABI; http://www3.appliedbiosystems.com/AB\_Home/index.htm) or primer sequences.

|  |  |  |  |
| --- | --- | --- | --- |
| **Gene symbol** **(alternative symbol)** | **Gene description** | **Main function\*\*** | **TaqMan assay or primer sequences** |
| *GAPDH***\*** | Glyceraldehyde 3-phosphate dehydrogenase | Housekeeping | Hs02786624\_g1 |
| *BMI1* | BMI1 polycomb ring finger oncogene | Maintains the transcriptionally repressive state of genes, involved in PR degeneration in mice and rats. | **F**: TTTTCCGGGATATTTTATCAAGCA**R**: GGGATTTAGCTCAGTGATCTTGATTC |
| *CCNA1***\*** | Cyclin A1 | Cell cycle and proliferation (controls and promotes G1/S (start) and G2/M (mitosis) transitions). | Cf02633425\_m1 |
| *CCNA2***\***(*CCNA/CCN1*) | Cyclin A2 | Cell cycle and proliferation (controls and promotes G1/S (start) and G2/M (mitosis) transitions). | Cf02695410\_m1 |
| *CCNB1***\*** | Сyclin B1; G2/mitotic-specific cyclin B1 | Cell cycle and proliferation (controls the G2/M transition). | **F**: GCCCTCTACCCCTGCATTTC**R**: TGCTCAACATCAACCTCTCCAA |
| *CCND1***\***(*BCL1/PRAD1*) | Cyclin D1; B-cell CLL/lymphoma 1 | Cell cycle and proliferation (regulatory subunit of CDK4 or CDK6, whose activity is required for G1/S transition). | **F**: CATCTACACTGACAACTCCATCC**R**: CAGGTTCCACTTCAGTTTGTTC |
| *CCND3* | Cyclin D3, G1/S-specific cyclin D3 | Cell cycle and proliferation (regulatory subunit of CDK4 or CDK6, whose activity is required for G1/S transition). | **F**: GTGTGTGCAGAGGGAGATCAAG**R**: CACAGACCTCCAGCATCCAGTA |
| *CCNE1***\*** | Cyclin E1 | Cell cycle and proliferation (G1/S-specific). | **F**: TGGCCCCAGTTTTTGCA**R**: CCTCTCGCAGTCCTGTCAATTT |
| *CDC25A* | Cell division cycle 25A | Phosphatase required for progression from G1 to S phase of the cell cycle, dephosphorylates CDK1 and thus activates it. | **F**: AACCTTGGCAATCGATGCA**R**: ACCGGGCGATGGAGCTA |
| *CDC25B* | Cell division cycle 25B | Phosphatase required for G2/M phases of the cell cycle progression, functions as a dosage-dependent inducer of mitotic progression, dephosphorylates CDK1 and thus activates it. | **F**: GAGCAGGCCATCCAAGCA**R**: GAAGCGTCGAATGGCAAACT |
| *CDC25C* | Cell division cycle 25C | Phosphatase required for progression of the cell cycle, functions as a dosage-dependent inducer of mitotic progression, when phosphorylated activates G2 cells into prophase, dephosphorylates CDK1 and thus activates it. | **F**: TCTGCCAACCTAAGCGTTTTG**R**: GATTCGAAAGATCAAGGCAACATT |
| *CDK1***\***(*CDC2*) | Cyclin-dependent kinase 1, Cell division control protein 2 homolog | Cell cycle and proliferation (essential for G1/S and G2/M transitions). | **F**: ACTCTTCAGAATTTTCAGAGCTTTGG**R**: GATTCCACTTCTGGCCACACTT |
| *CDK2* | Cyclin-dependent kinase 2 | Cell cycle and proliferation (activity restricted and essential for G1/S transition, involved in PR degeneration in mice and rats). | **F**: CCTCATCAAGAGCTATCTGTTCCA**R**: CCCGATGAGAATGGCAGAAG |
| *CDK4***\*** | Cyclin-dependent kinase 4 | Cell cycle and proliferation (activity restricted and essential for G1/S transition, phosphorylates RB1, involved in PR degeneration in mice and rats). | **F**: CAGTGGAGACCATCAAGGATCTG**R**: GCAGTTGGCATGAAGGAAATCT |
| *CDK6***\*** | Cyclin-dependent kinase 6 | Cell cycle and proliferation (essential for G1/S transition, involved in PR degeneration in mice and rats. | **F**: CAGTGGTCGTCACACTGTGGTA**R**: GGTGGCATAGCTGGACTGAAG |
| *CDKN1A* (*P21, CIP1*) | Cyclin-dependent kinase inhibitor 1A | Potent cyclin-dependent kinase inhibitor, binds to and inhibits activity of cyclin-CDK2 or -CDK4 complexes, regulator of cell cycle progression at G1. | **F**: CACCTCTCAGGGCCGAAAA**R**: GGCGTTTGGAGTGATAGAAATCTG |
| *CDKN1B* (*P27, KIP1*) | Cyclin-dependent kinase inhibitor 1B | Cyclin-dependent kinase inhibitor, binds to and prevents activation of CCNE-CDK2 or CCND-CDK4 complexes, controls cell cycle progression at G1. | **F**: CCGACGATTCCTCTCCTCAA**R**: GGAACCGTCTGAGACGTTTTCT |
| *CDKN2A* (*P16*, *INK4A*) | Cyclin-dependent kinase inhibitor 2A | Cyclin-dependent kinase inhibitor, induces cell cycle arrest in G1 and G2 phases, acts as a tumor suppressor, induces G2 arrest and apoptosis in a p53-independent manner by preventing activation of cyclin B1/CDC2 complexes. | **F**: TACGGAAGGTGCGGAAGGTC**R**: TGAAAAAGGAGTGCTCTGGGC |
| *CRB1* | Crumbs homolog 1 (Drosophila) | Localizes to the inner segment of mammalian PRs, plays a role in PR morphogenesis. | **F**: CCTACGAAGGCCCGAACTG**R**: AACTCTTGTCAAGATTAAAAGCAGCAA |
| *E2F1***\*** | E2F transcription factor 1, Retinoblastoma-associated protein 1 | Cell cycle and proliferation (can mediate both cell proliferation and p53-dependent/independent apoptosis, involved in PR degeneration in mice and rats. | **F**: CTCCAAGCCATAGACTCCTCAGA**R**: GGGCACAGGAAAACGTCAAT |
| *GRK1* (*RHOK/GPRK1/RK*) | G protein-coupled receptor kinase 1, rhodopsin kinase | Phosphorylates rhodopsin to initiate its deactivation. | **F**: AGGGTGCTATGGTGGAGAAGAA**R**: GTTTCAAATGCATAGGCCAGAGA |
| *LATS1***\***(*WARTS/wts*) | LATS, large tumor suppressor, homolog 1 (Drosophila) | Hippo pathway, tumor suppressor gene. | Cf02626754\_m1 |
| *LATS2***\*** | LATS, large tumor suppressor, homolog 2 (Drosophila) | Hippo pathway, tumor suppressor gene. | Cf02633924\_m1 |
| *MOB1A***\***(*MOBKL1A/MOB1B/MATS2*) | MOB kinase activator 1A | Hippo pathway, activates LATS1 and LATS2. | Cf02649192\_m1 |
| *NDR1***\***(*STK38/NDR*) | Serine/threonine kinase 38 | Regulates cell cycle, proliferation, apoptosis. | **F**: CAACCTTCTCCTGGACAGCAA**R**: CTGTGCAAAGGCCGAAGTC |
| *NRL* (*RP27)* | Neural retina leucine zipper | Regulates PR development  | Cf02655725\_m1 |
| *NR2E3* (*ESCS/PNR/RNR/rd7*) | Nuclear receptor subfamily 2, group E, member 3 | Proliferation; activates rod and represses cone development. | **F**: TGAAGGCCCTGGTCCTCTT**R**: ACGTGCTCAGGATCCTTCAG**TaqMan:** CACACCAGAAACTCG |
| *PAX6***\***(*AN/AN2/MGDA/WAGR*) | Paired homeobox gene 6 | Regulates eye development. | Cf02649657\_g1 |
| *PCNA* | Proliferating cell nuclear antigen | Cell proliferation. | Cf02634115\_m1 |
| *RB1***\*** | retinoblastoma 1, osteosarcoma | Negative regulator of cell cycle and proliferation. | **F**: TTATCGAGTAATGGAATCCATGCTTA**R**: TGTCGTCATTCAGGAGTTTGCT |
| *RBP3* (*IRBP*) | Retinol binding protein 3, interstitial, interphotoreceptor retinoid-binding protein | Shuttles retinoids in PRs. | Cf02676369\_m1 |
| *RCVRN* (*RCV1*) | Recoverin, cancer-associated retinopathy protein | Inhibits rhodopsin kinase, regulates sensory adaptation in retina. | Cf02642703\_m1 |
| *RDS* (*PRPH2/CACD2/DS/RP7/rd2*) | Peripherin 2 (retinal degeneration, slow) | Prevents PR degeneration. | Cf02628333\_m1 |
| *STK38L***\***(*NRD2*) | Serine/threonine kinase 38 like | Regulates cell cycle, proliferation, and apoptosis. | Cf02709228\_m1 (spans exons 4-5)Cf02634613\_m1 (spans exon 6-7) |

**\***: subset of genes tested in RPE.

**\*\***: gene function information mainly based on GeneCards ([www.genecards.org](Supplemental%20Table%20S2%20genes-finale.docx)).