**Figure 5—source data 4: Cone mRNA raw reads from RNA-seq of all 35 retinas used in the study.**

|  |  |
| --- | --- |
| **Gene** | **Raw reads** |
| ***Slc2a1*** | 278.8 ± 21.4  |
| ***Ldha*** | 6827.4 ± 511.8  |
| ***Ldhb*** | 296.0 ± 28.7  |

*Data presented as: Mean ± SEM (n=35 for all).*

*No significant difference in these gene with or without Txnip in each strain.*