|  |  |  |
| --- | --- | --- |
| **Figure 4B** | P value | Summary |
| ANOVA | 0.0141 | \* |
| Šídák's multiple comparisons test | Adjusted P Value | Summary |
| STOP siCTRL vs. STOP siTPR | 0.6882 | ns |
| STOP siCTRL vs. RAS siCTRL | 0.6959 | ns |
| RAS siCTRL vs. RAS siTPR | 0.0476 | \* |

|  |  |  |
| --- | --- | --- |
| **Figure 4D** | P value | Summary |
| ANOVA | 0.0108 | \* |
| Šídák's multiple comparisons test | Adjusted P Value | Summary |
| STOP siCTRL vs. STOP siTPR | 0.9333 | ns |
| STOP siCTRL vs. RAS siCTRL | 0.0109 | \* |
| STOP siCTRL vs. RAS siTPR | 0.1124 | ns |
| RAS siCTRL vs. RAS siTPR | 0.6315 | ns |
| STOP siTPR vs RAS siTPR | 0.1734 | ns |

**Figure 4 – source data 1. Statistical analysis for *STING1* qPCR data in Figure 4B and for cGAMP ELISA data in Figure 4D.** One-way ANOVA was used to determine statistical significance followed by Šídák's multiple comparisons test.