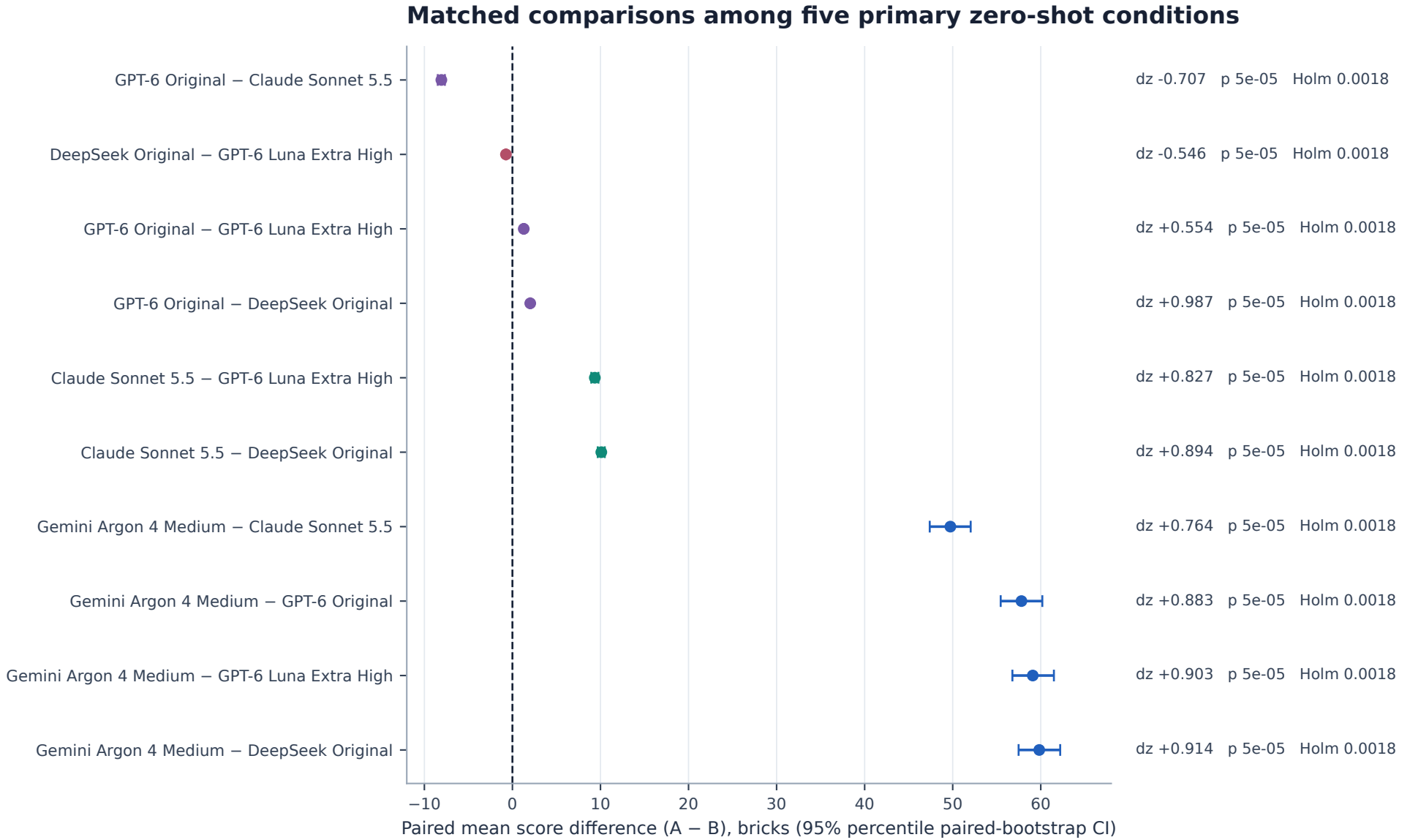


Source: corrected Phase 2 paired-analysis JSON · 3,000 matched pairs per row. P-values are two-sided; Holm adjustment uses the full 36-pair family. Nonsignificance is not evidence of equivalence.