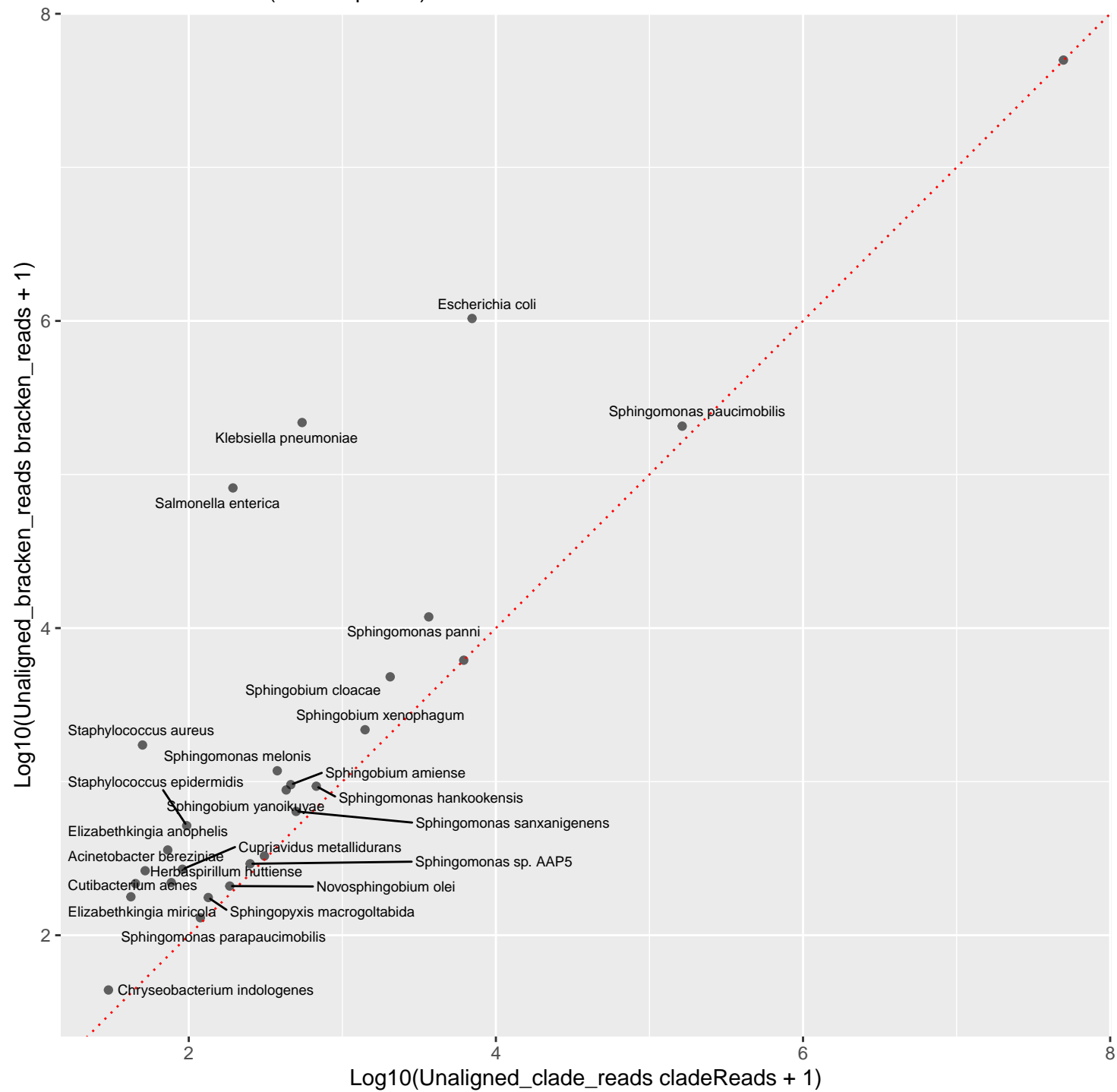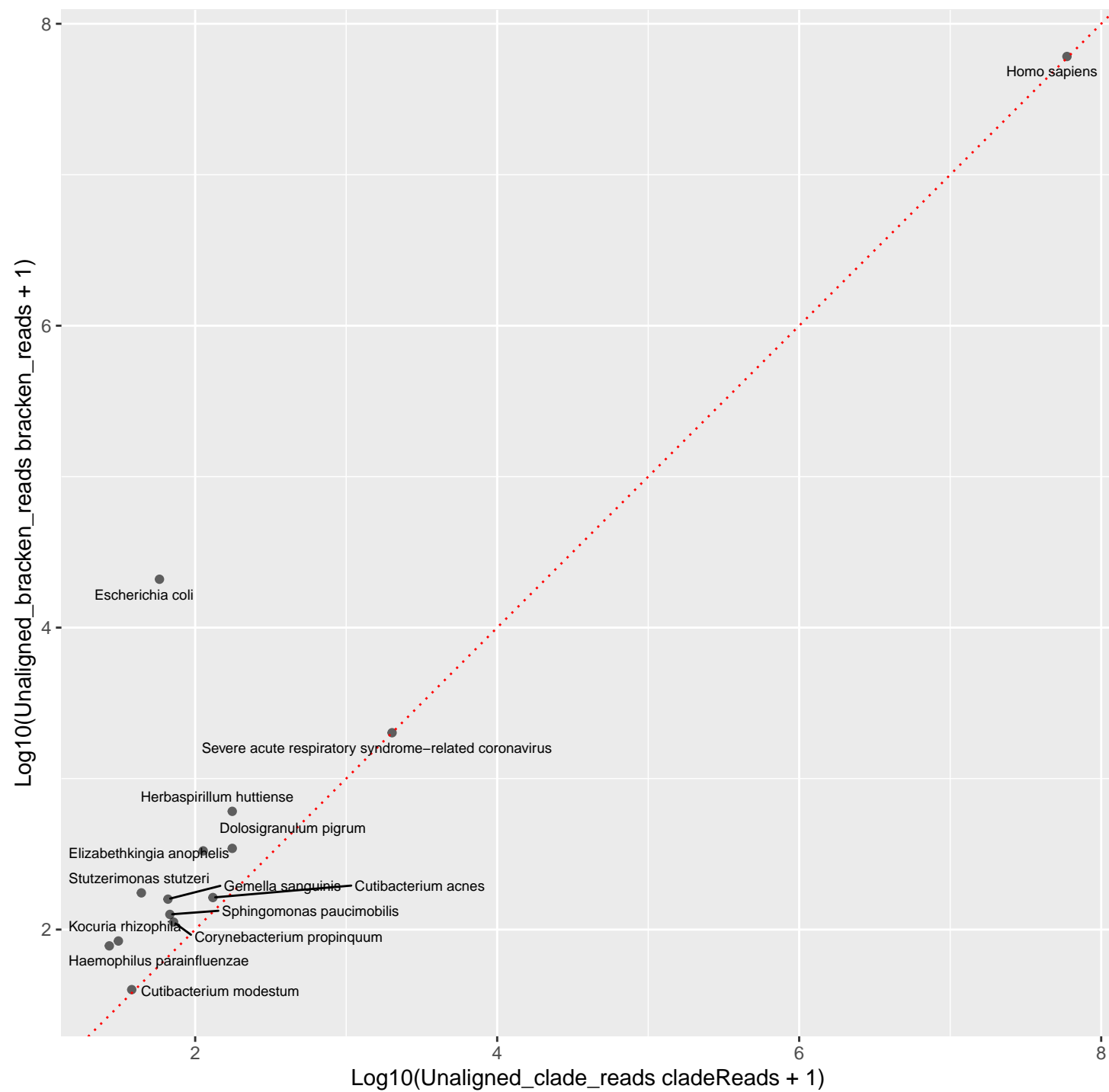Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (Neg_Control_W)

Pearson r = 0.84726 (n = 28 species)

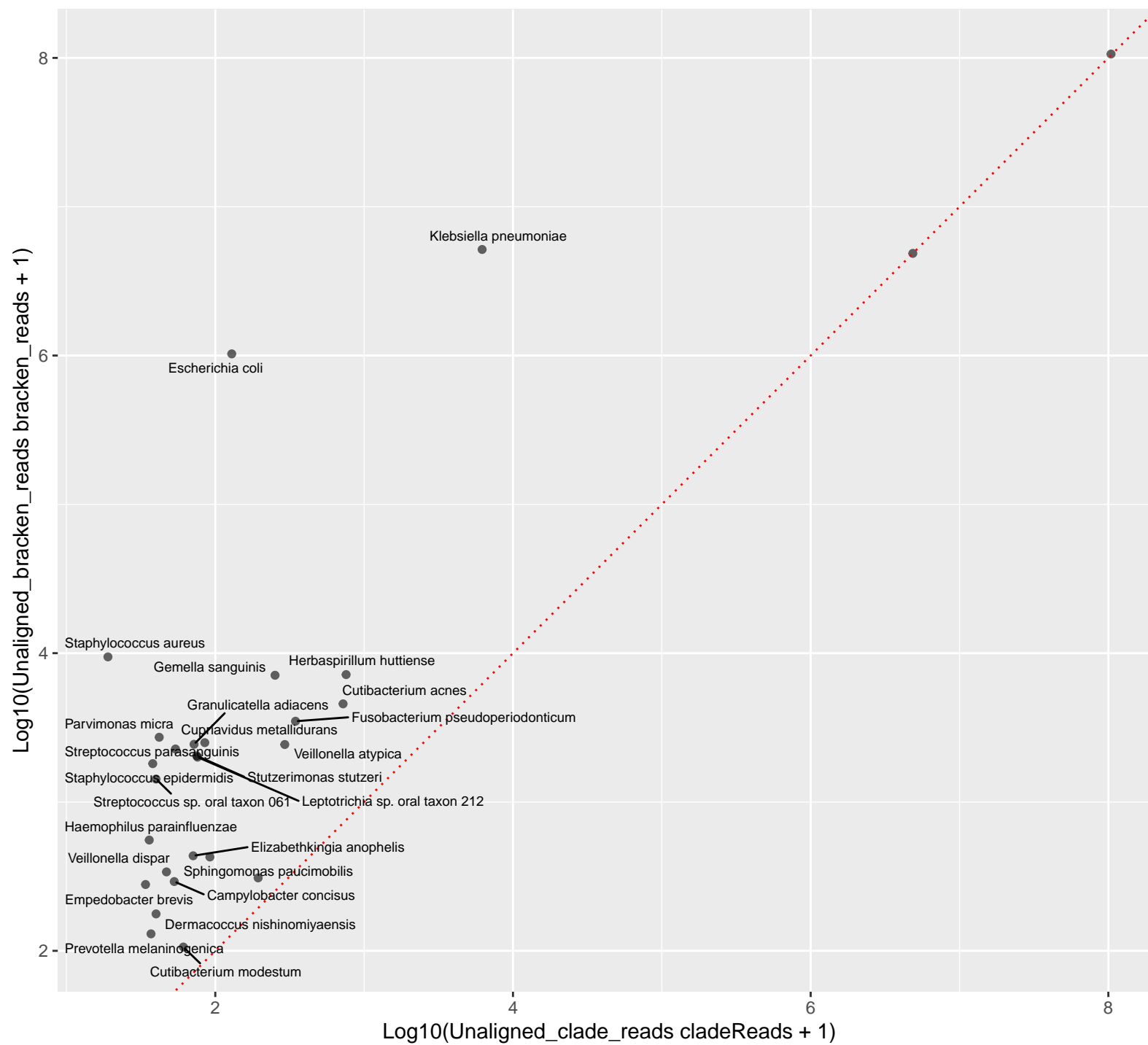**Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W61303224)**
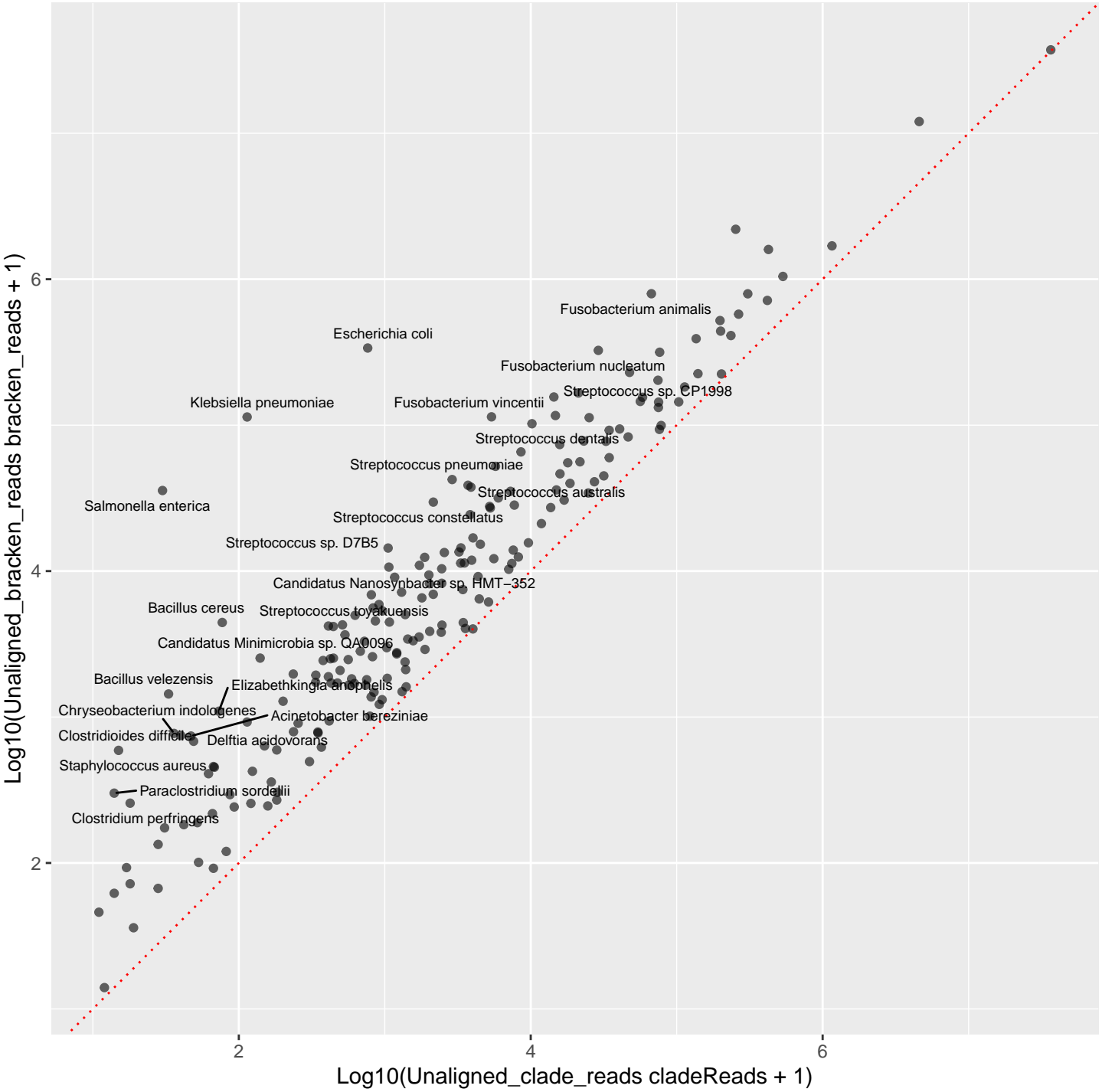
Pearson r = 0.92102 (n = 14 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W61304616)
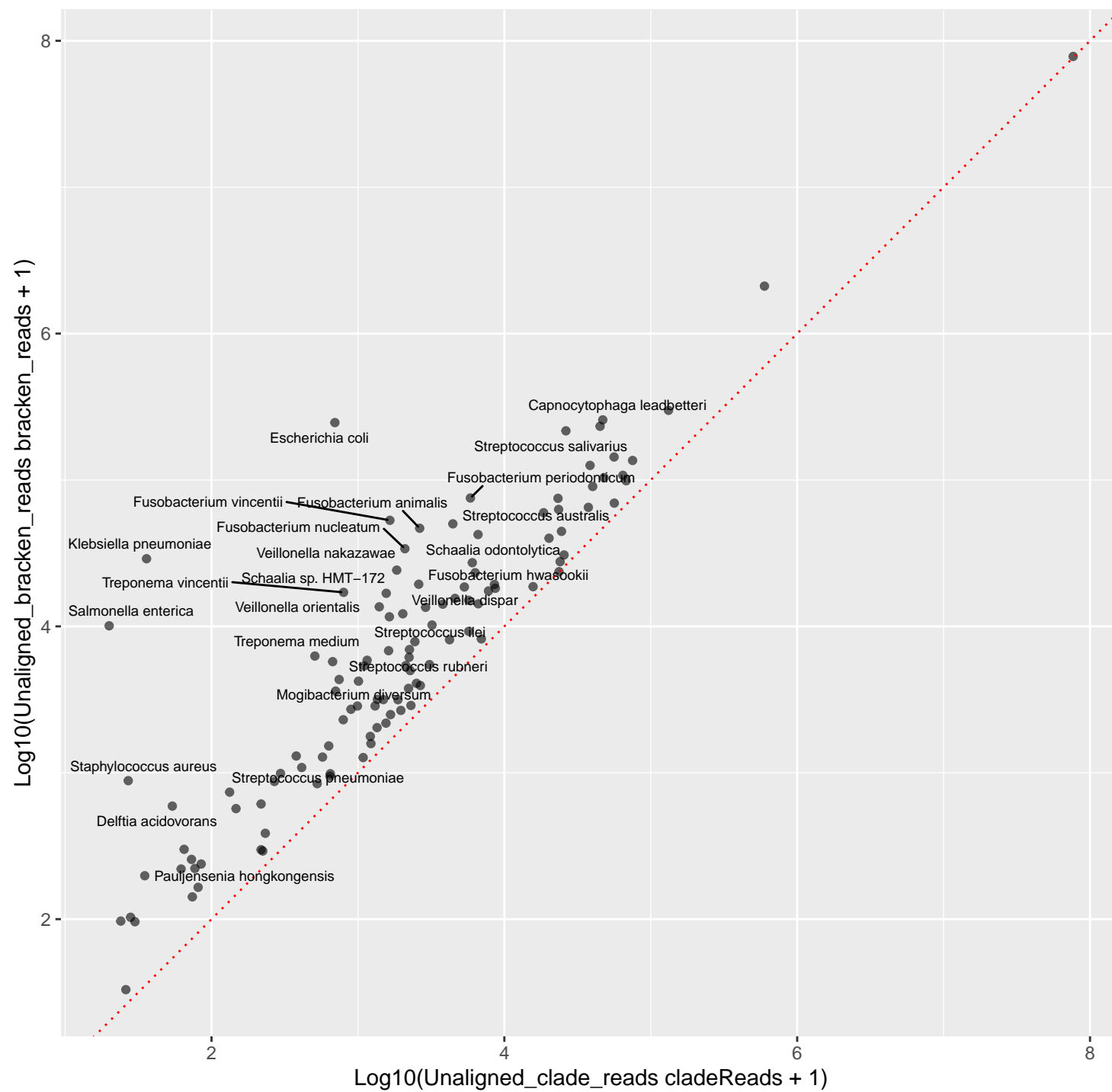
Pearson r = 0.8318 (n = 28 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W61402281)
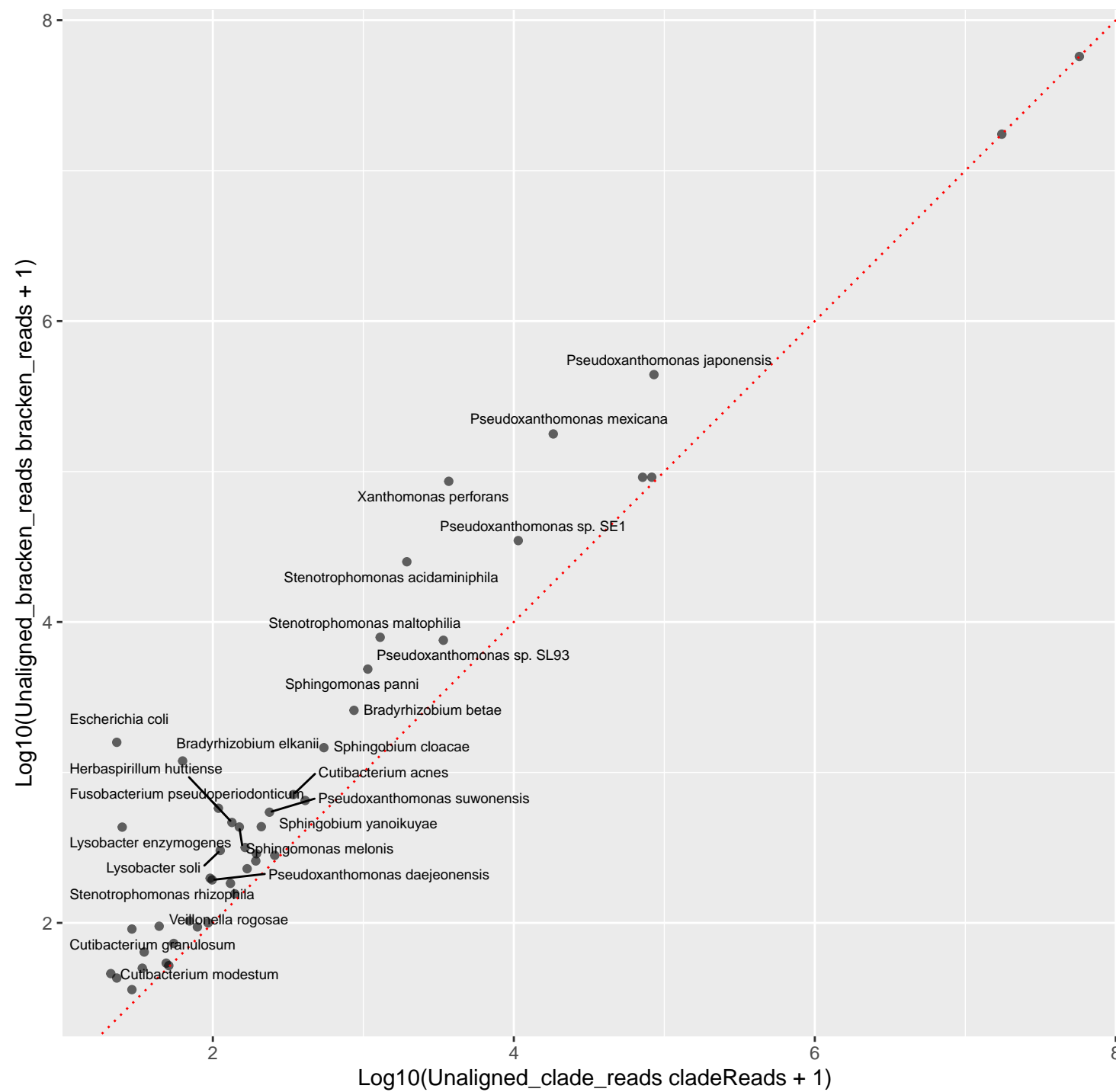
Pearson r = 0.92672 (n = 199 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W61402322)
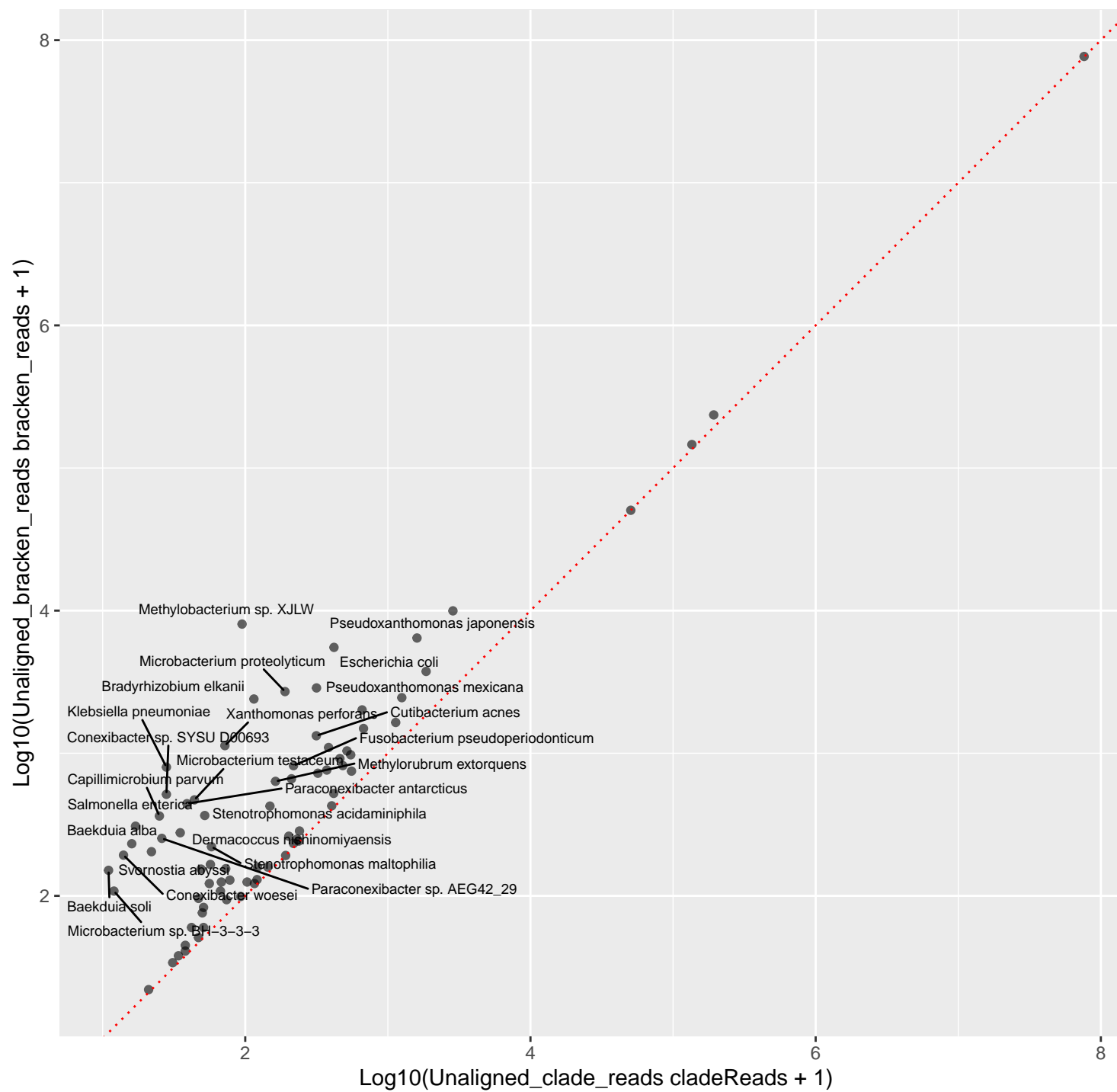
Pearson r = 0.88894 (n = 116 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W61501463)
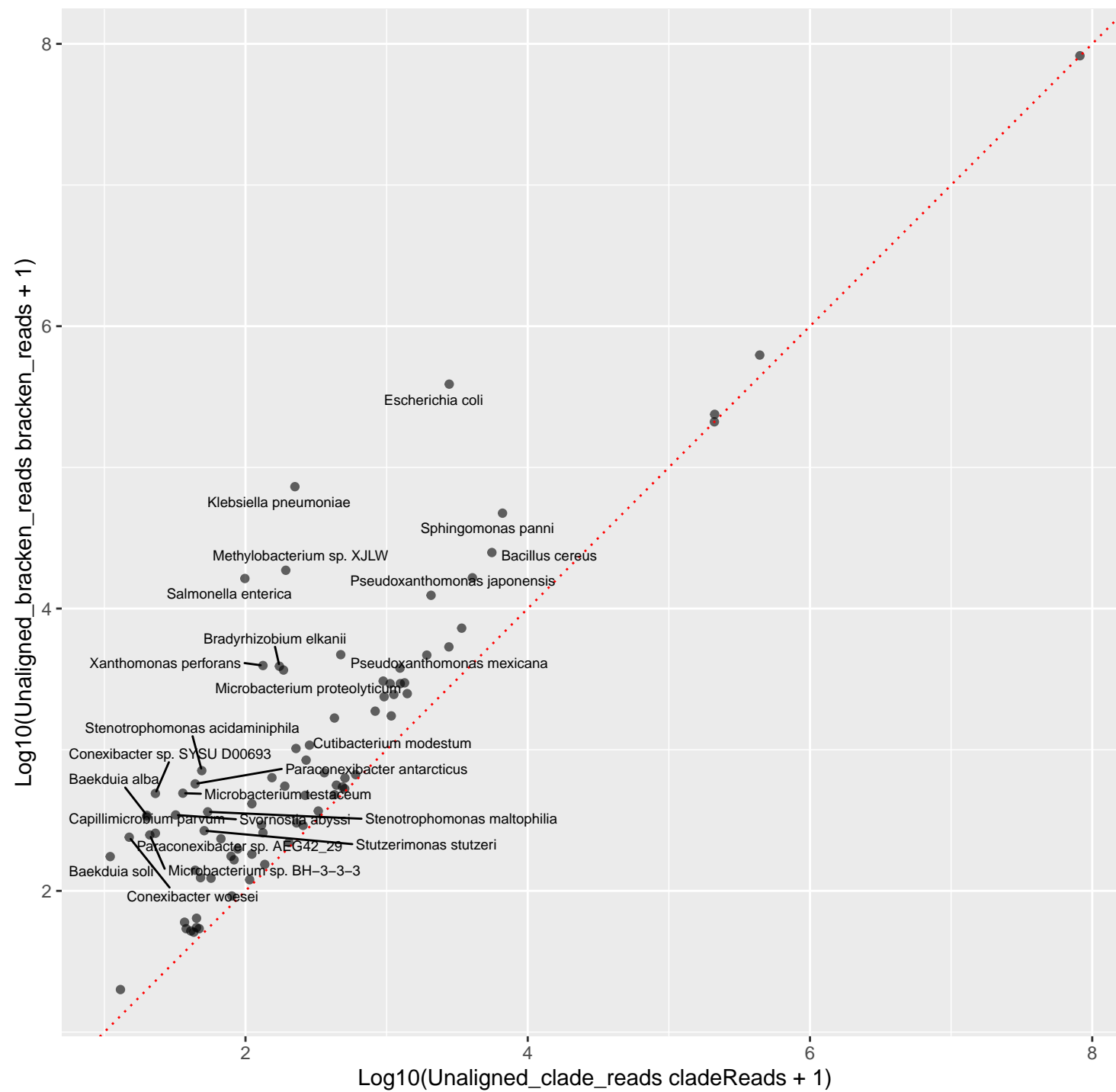
Pearson r = 0.95804 (n = 47 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W71607201)

Pearson r = 0.89705 (n = 78 species)

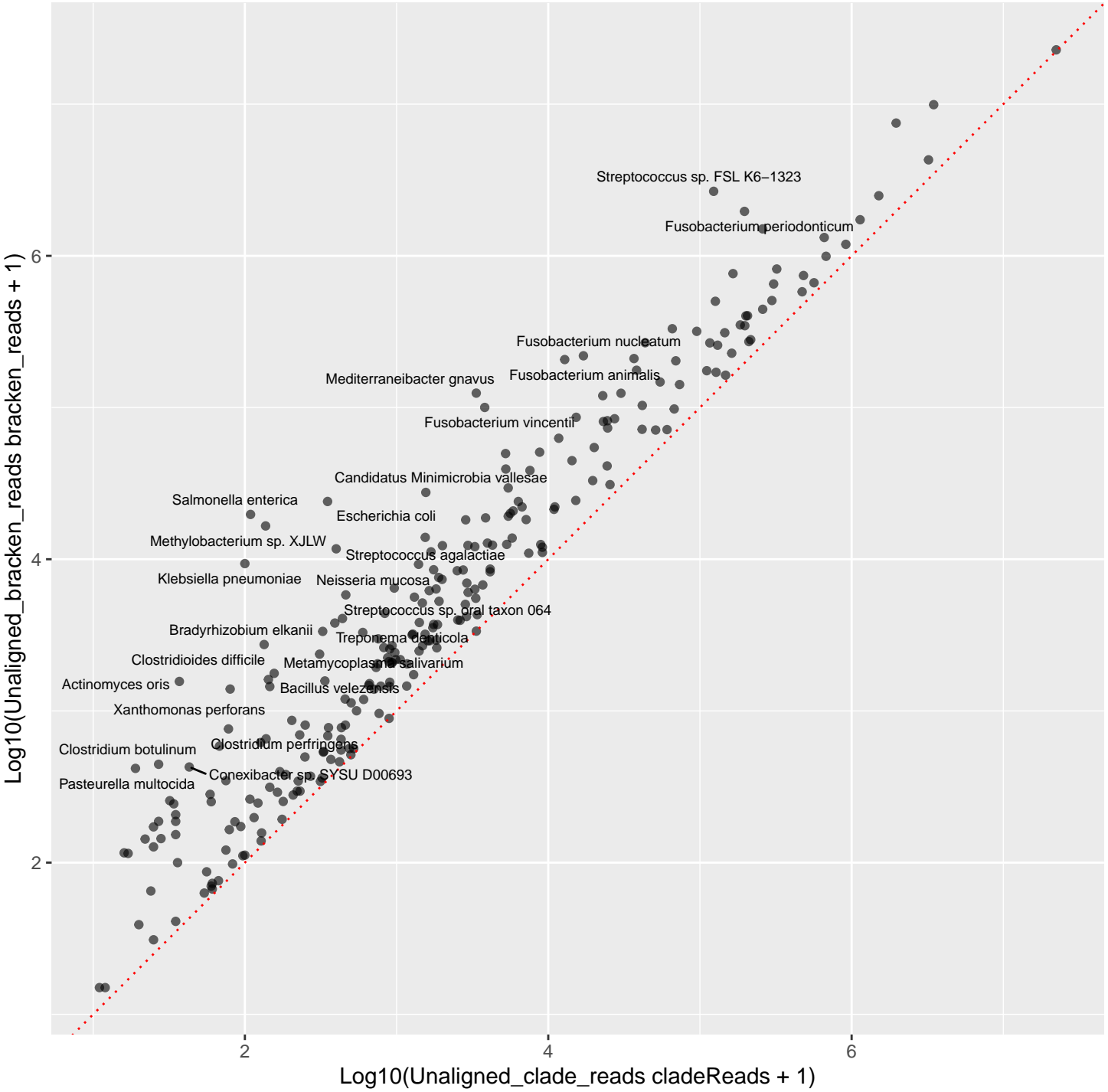Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W72504187)
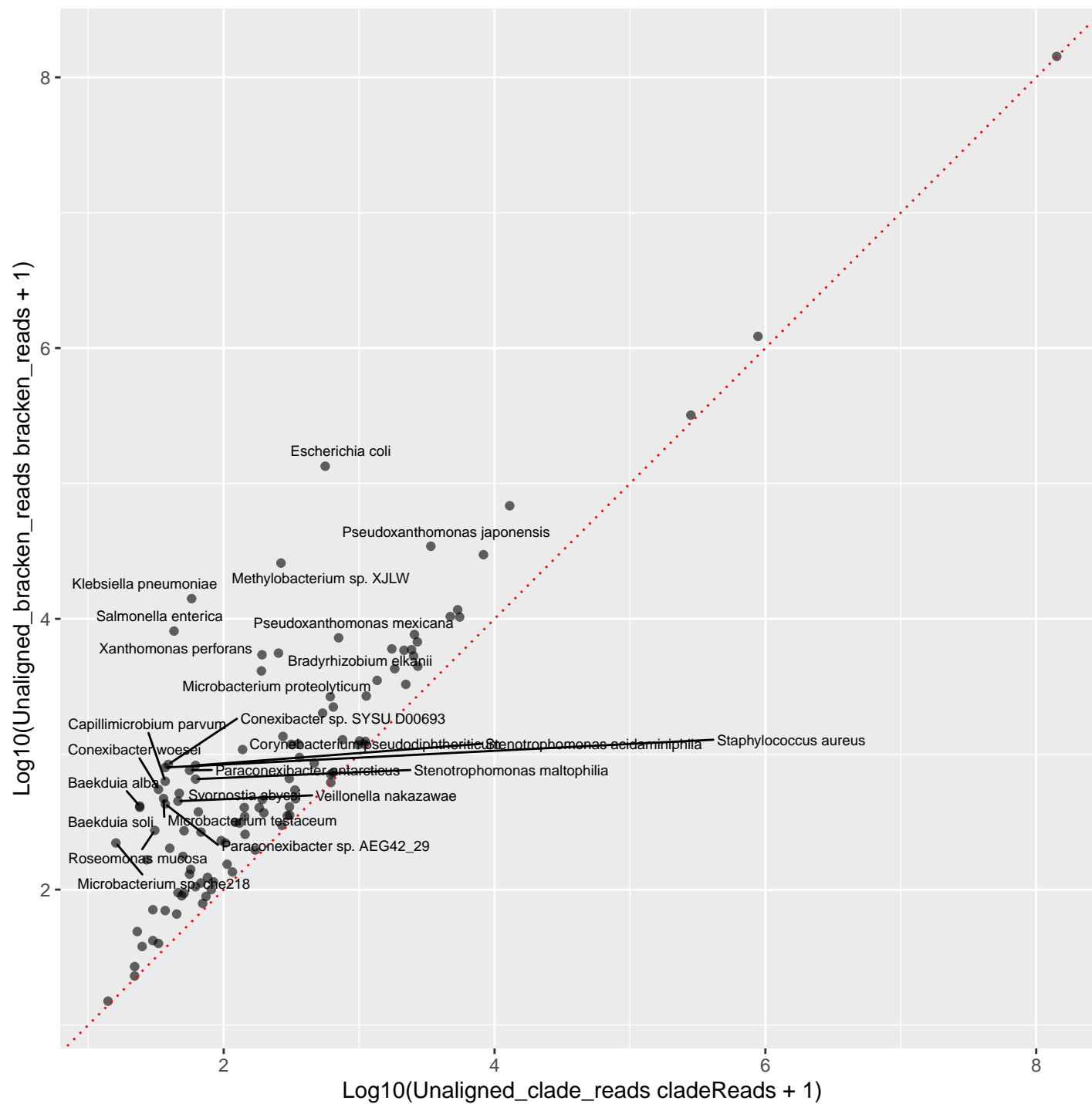
Pearson r = 0.87096 (n = 82 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W72803699)
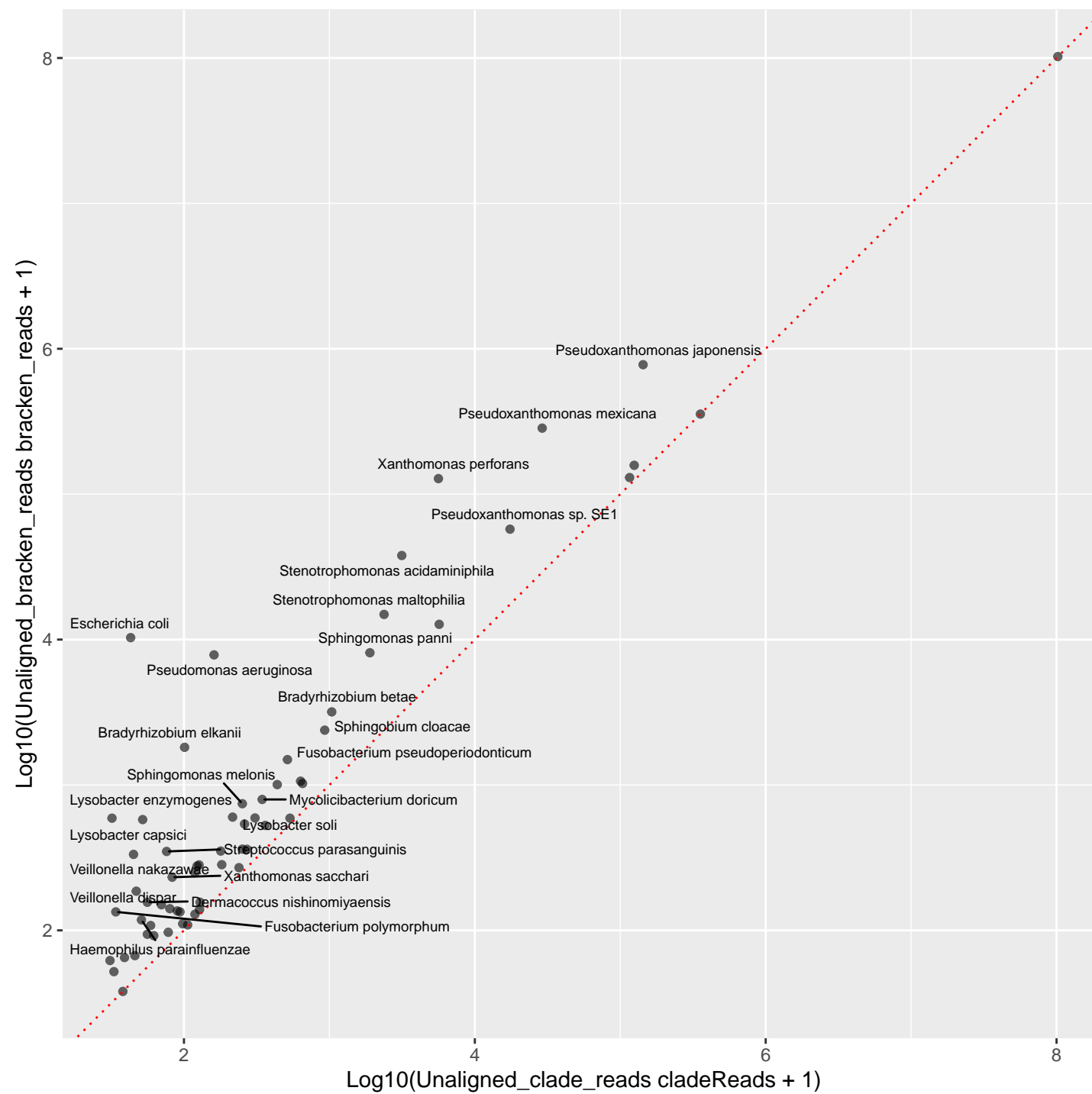
Pearson r = 0.95301 (n = 256 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W72902029)
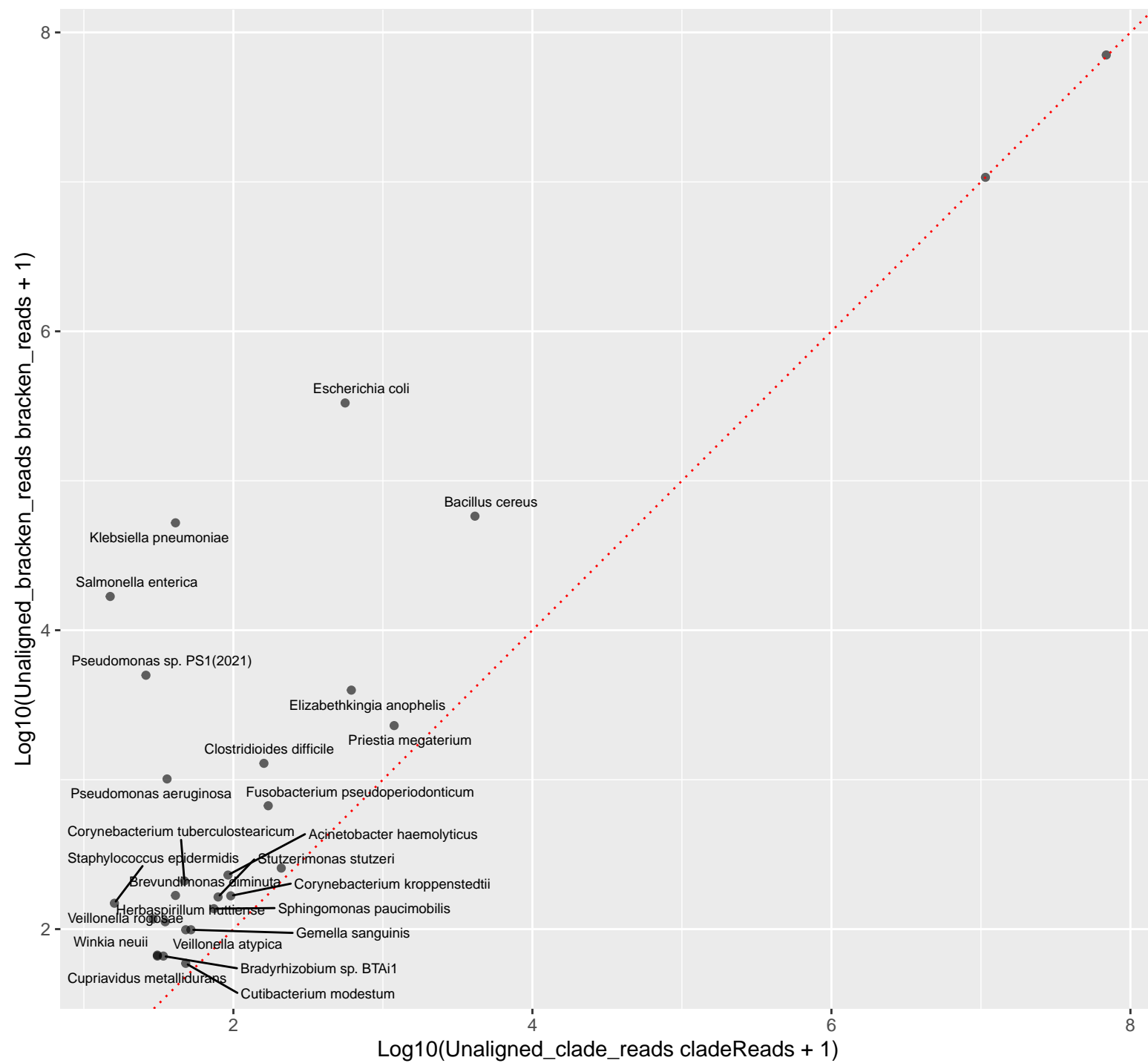
Pearson r = 0.87012 (n = 108 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W73007482)
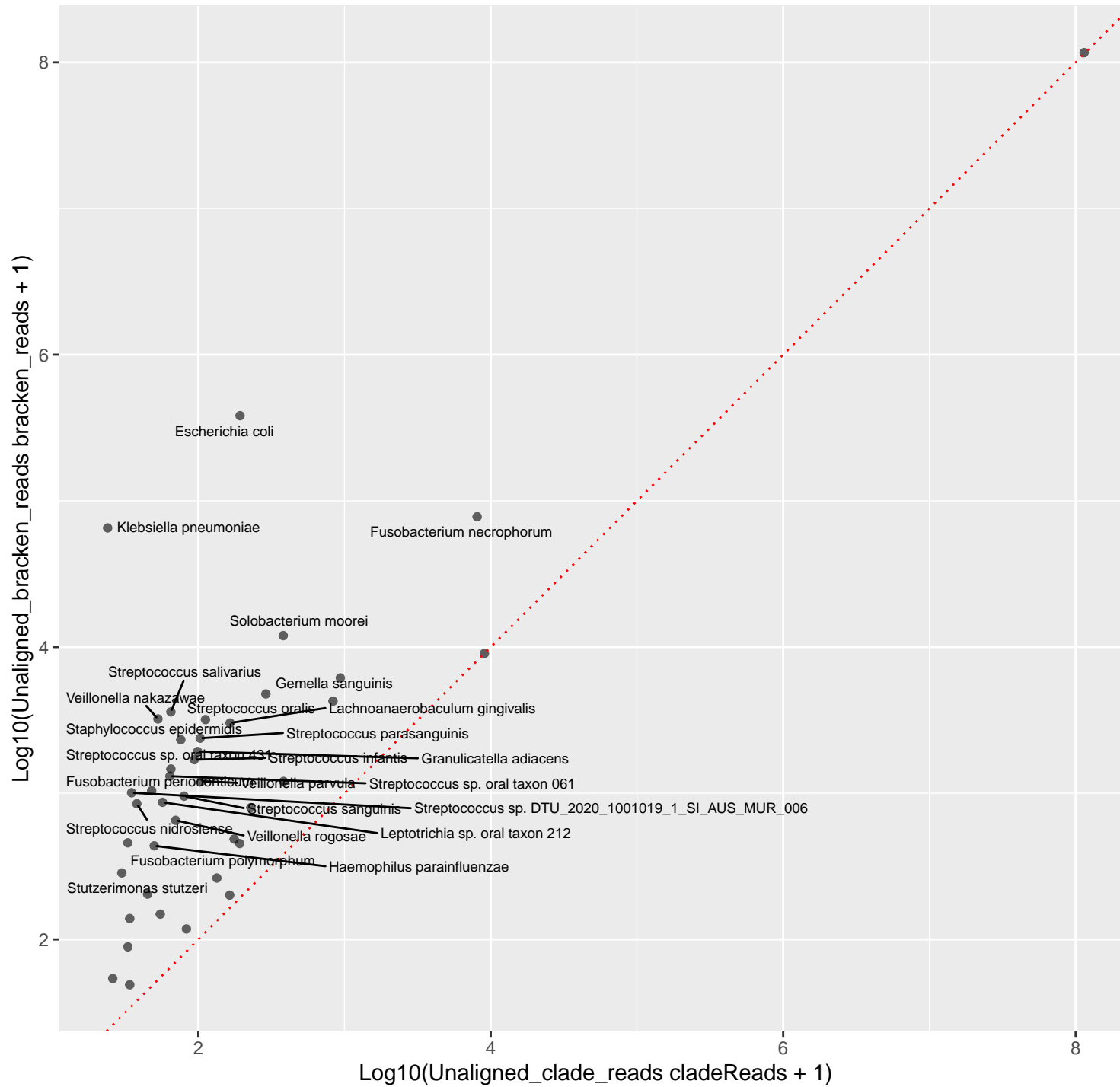
Pearson r = 0.93448 (n = 63 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W73007677)
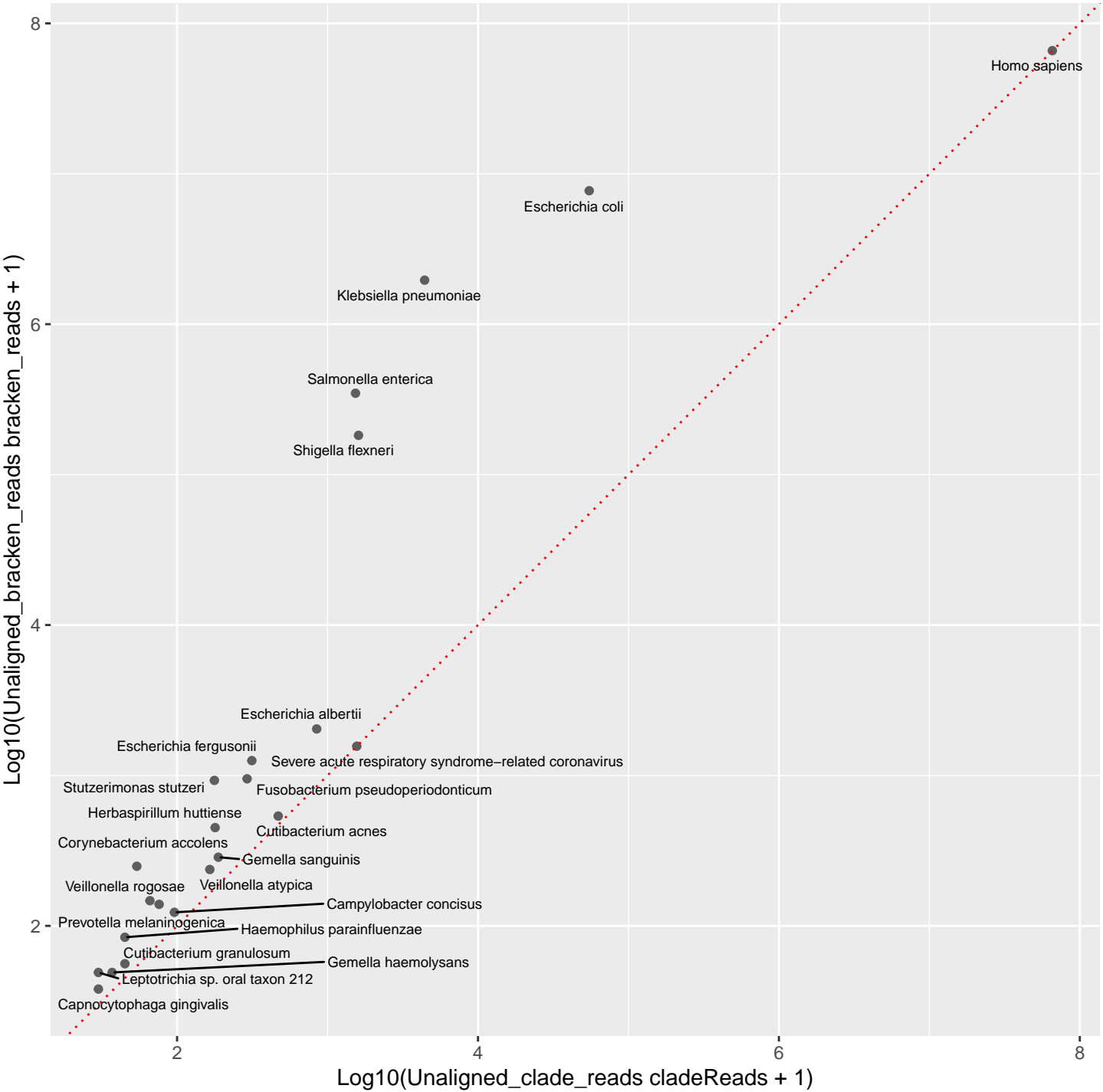
Pearson r = 0.83598 (n = 28 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W73007934)
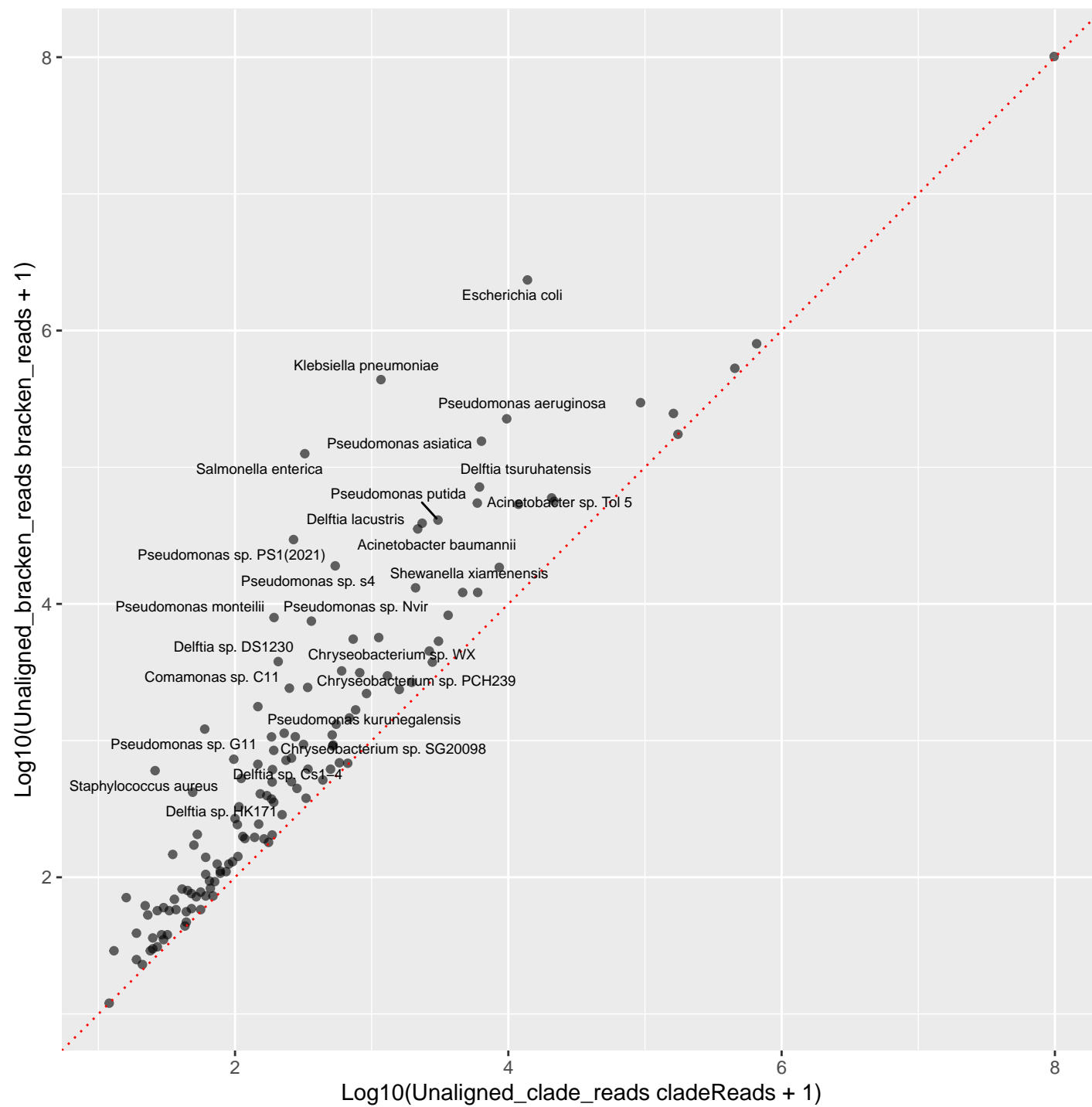
Pearson r = 0.7845 (n = 42 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W73109775)
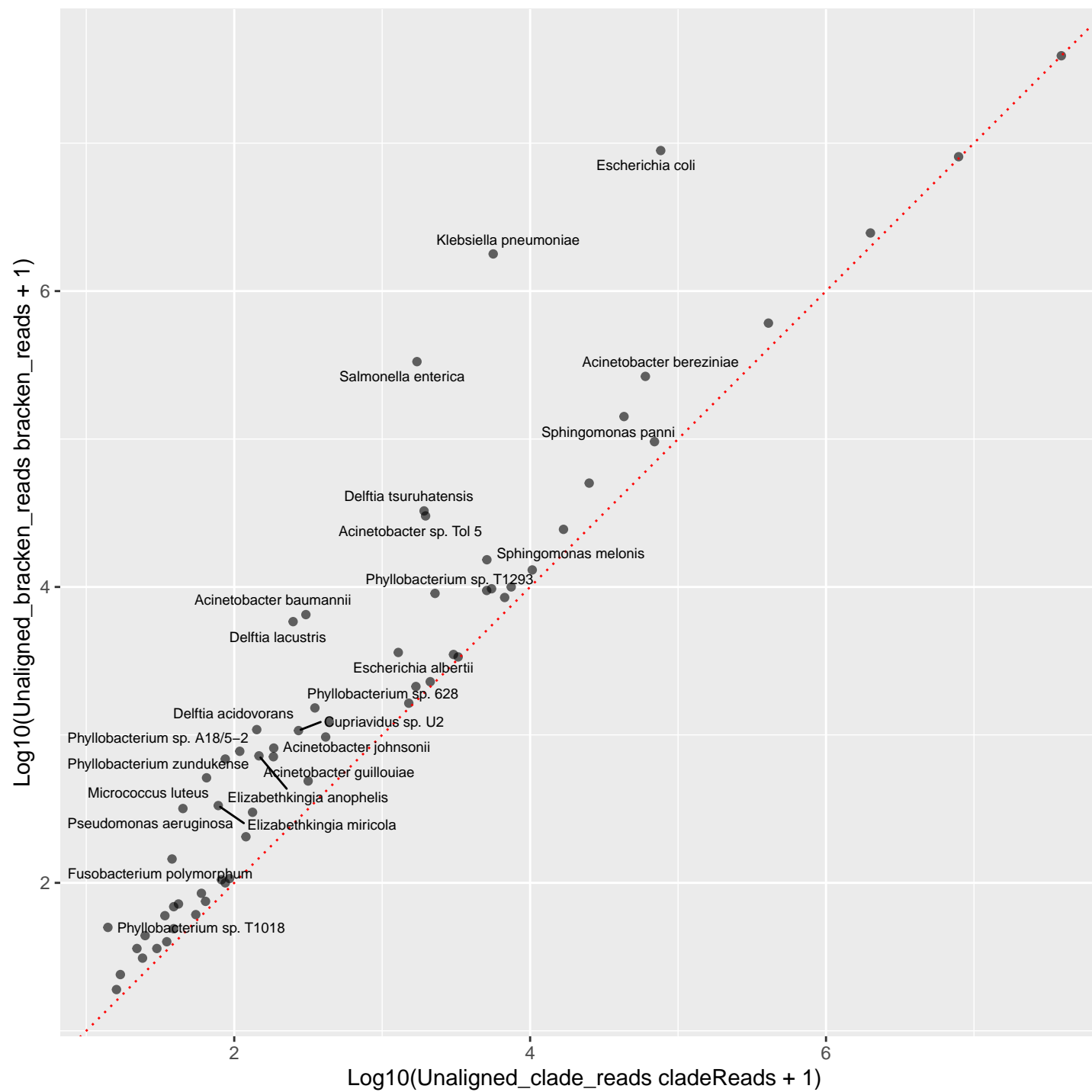
Pearson r = 0.89996 (n = 23 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W81802779)

Pearson r = 0.91646 (n = 134 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W90303891)
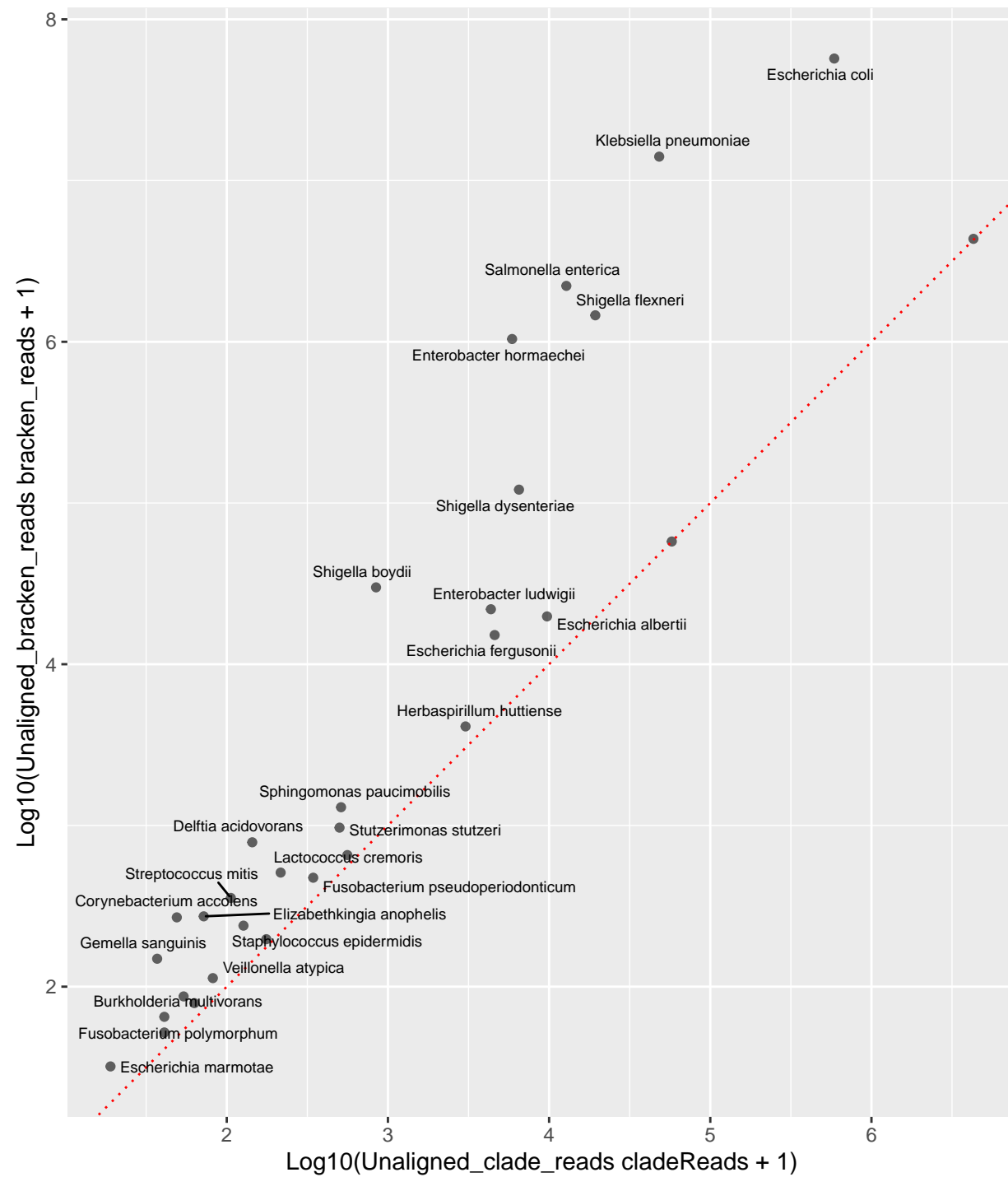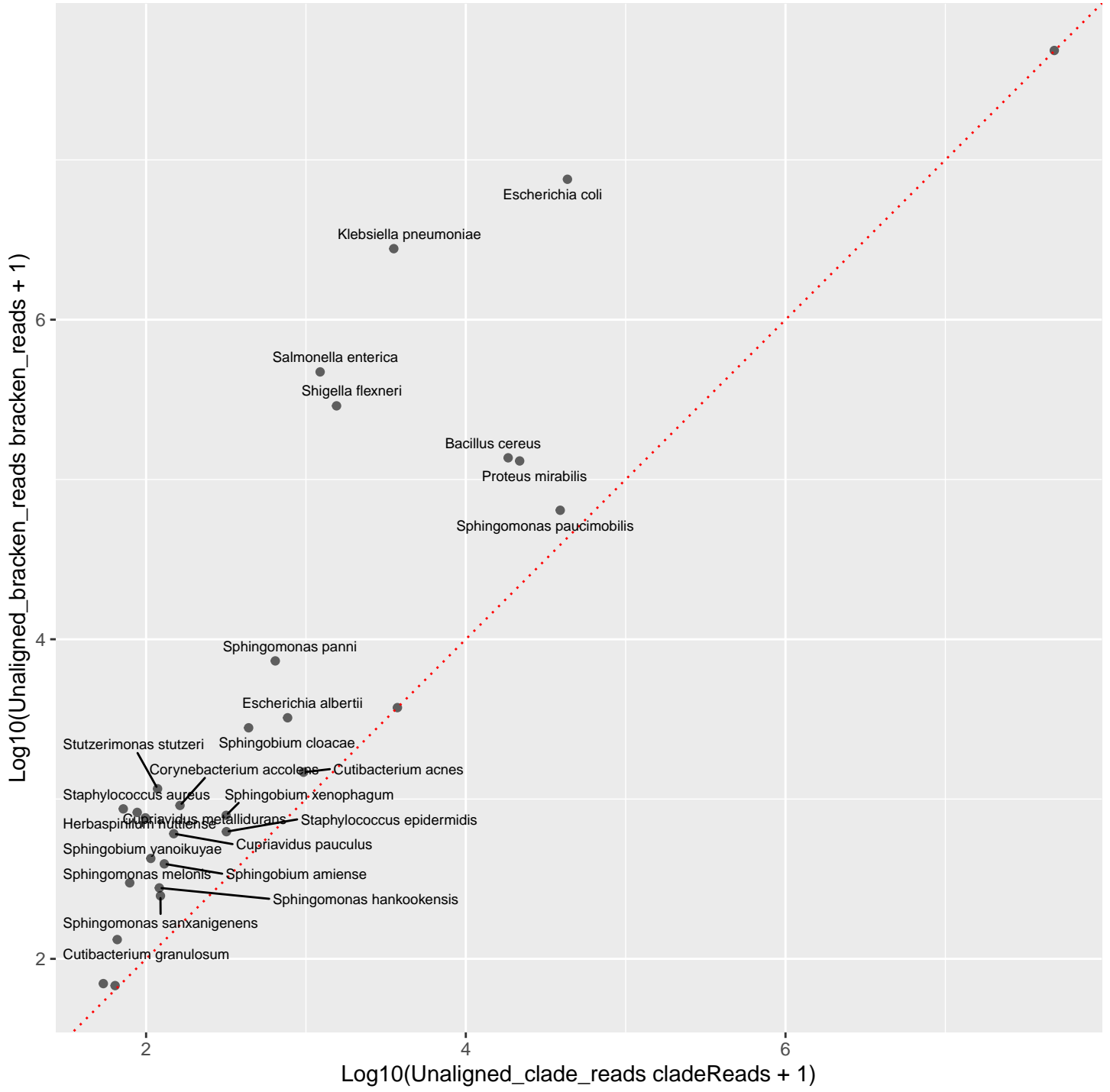
Pearson r = 0.93633 (n = 64 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (W90504037)
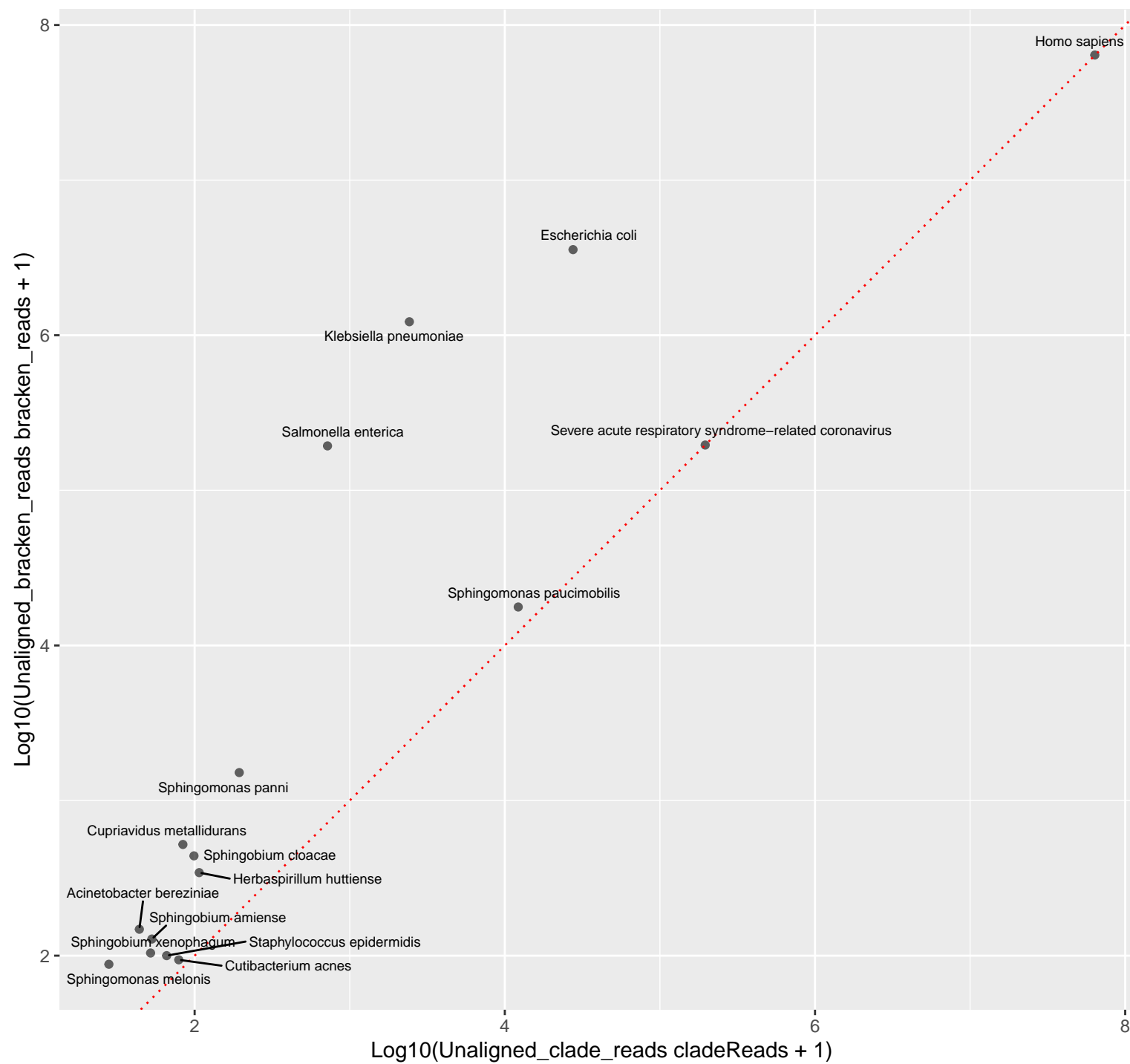
Pearson r = 0.92638 (n = 31 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (X01703004)

Pearson r = 0.87163 (n = 29 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (X22106916)

Pearson r = 0.89446 (n = 16 species)

Correlation: Unaligned_clade_reads vs Unaligned_bracken_reads (X30306045)

Pearson r = 0.90344 (n = 226 species)