

Figure S1a. Q-Q plots (for the models evaluating the **mesiodistal crown diameter**) for the best fit regression models to assess if the set of residuals are normally distributed for the maxillary (A–E) and mandibular (G–J) dentoalveolar structures.

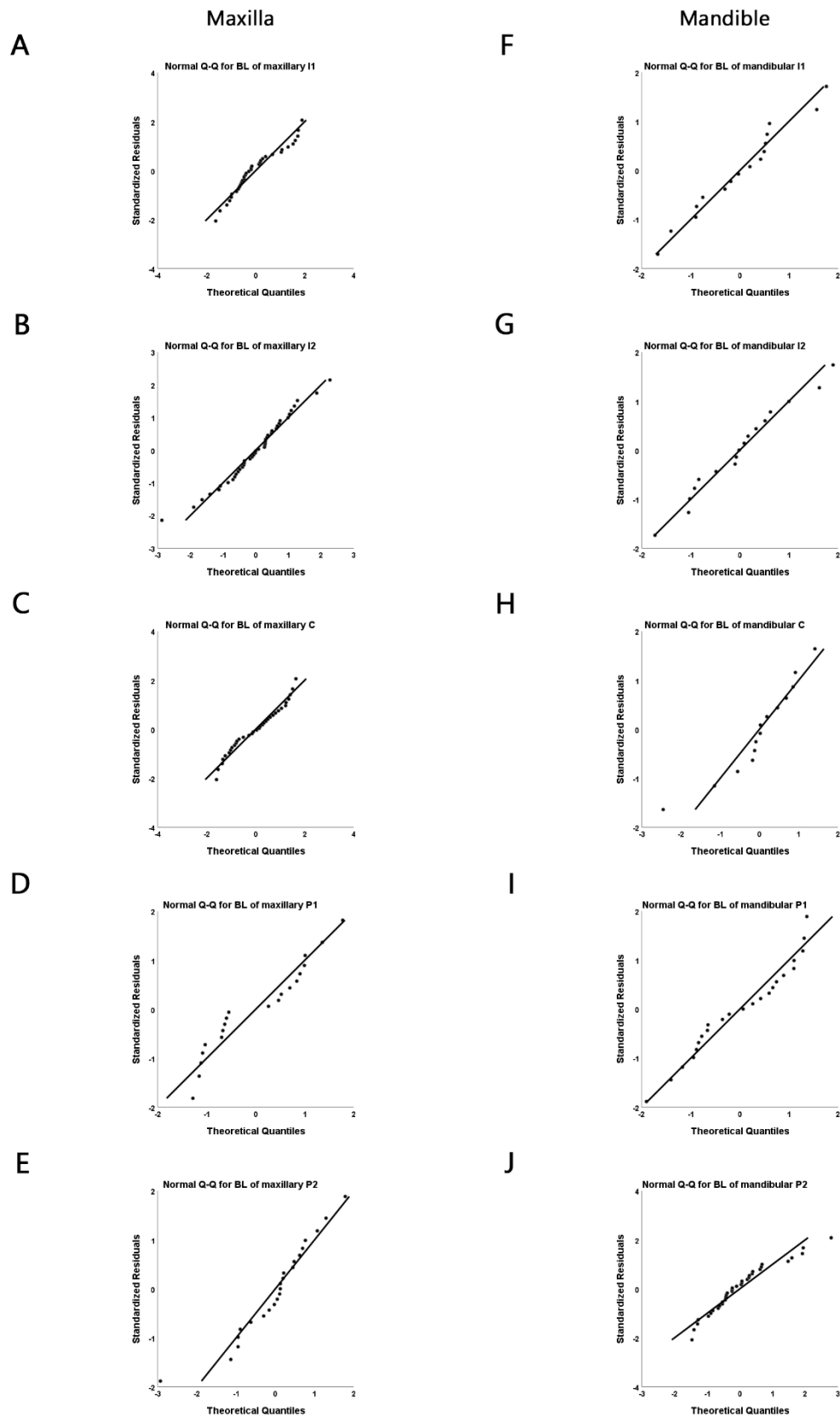


Figure S1b. Q-Q plots (for the models evaluating the **buccolingual crown diameter**) for the best fit regression models to assess if the set of residuals are normally distributed for the maxillary (A–E) and mandibular (G–J) dentoalveolar structures.

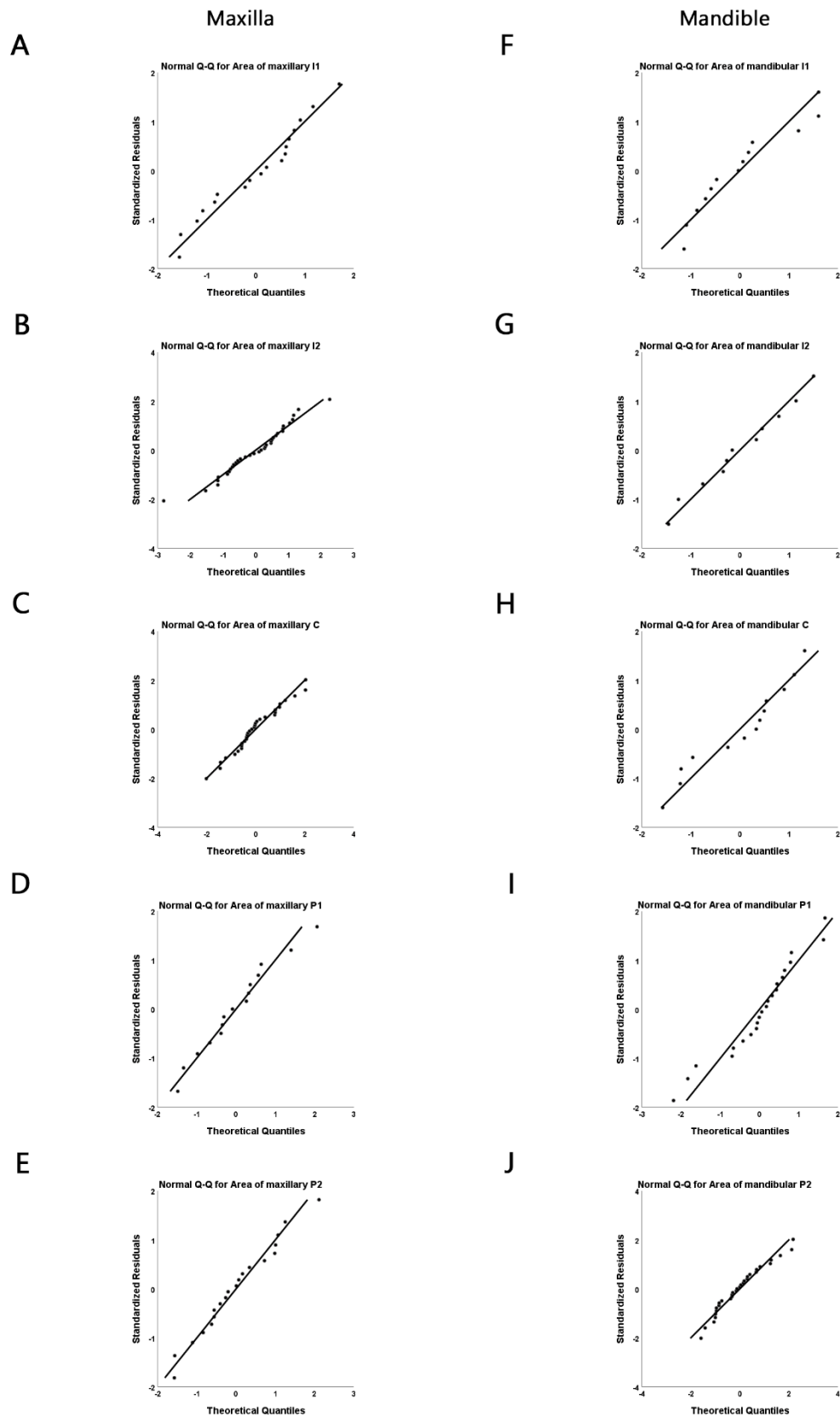


Figure S1c. Q-Q plots (for the models evaluating the **crown area**) for the best fit regression models to assess if the set of residuals are normally distributed for the maxillary (A–E) and mandibular (G–J) dentoalveolar structures.

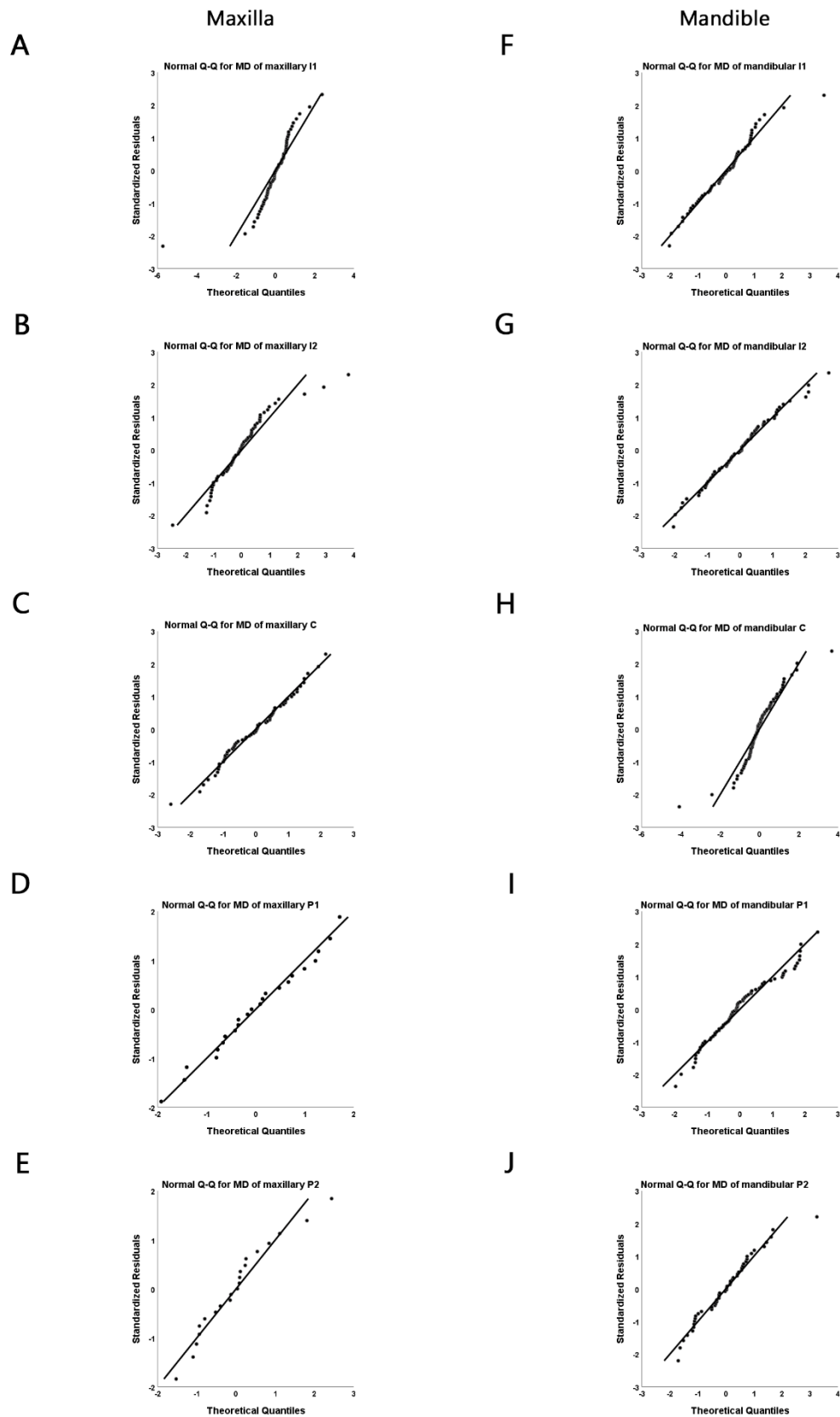


Figure S1d. Q-Q plots (for the models evaluating the **mesiodistal cervical diameter**) for the best fit regression models to assess if the set of residuals are normally distributed for the maxillary (A-E) and mandibular (G-J) dentoalveolar structures.

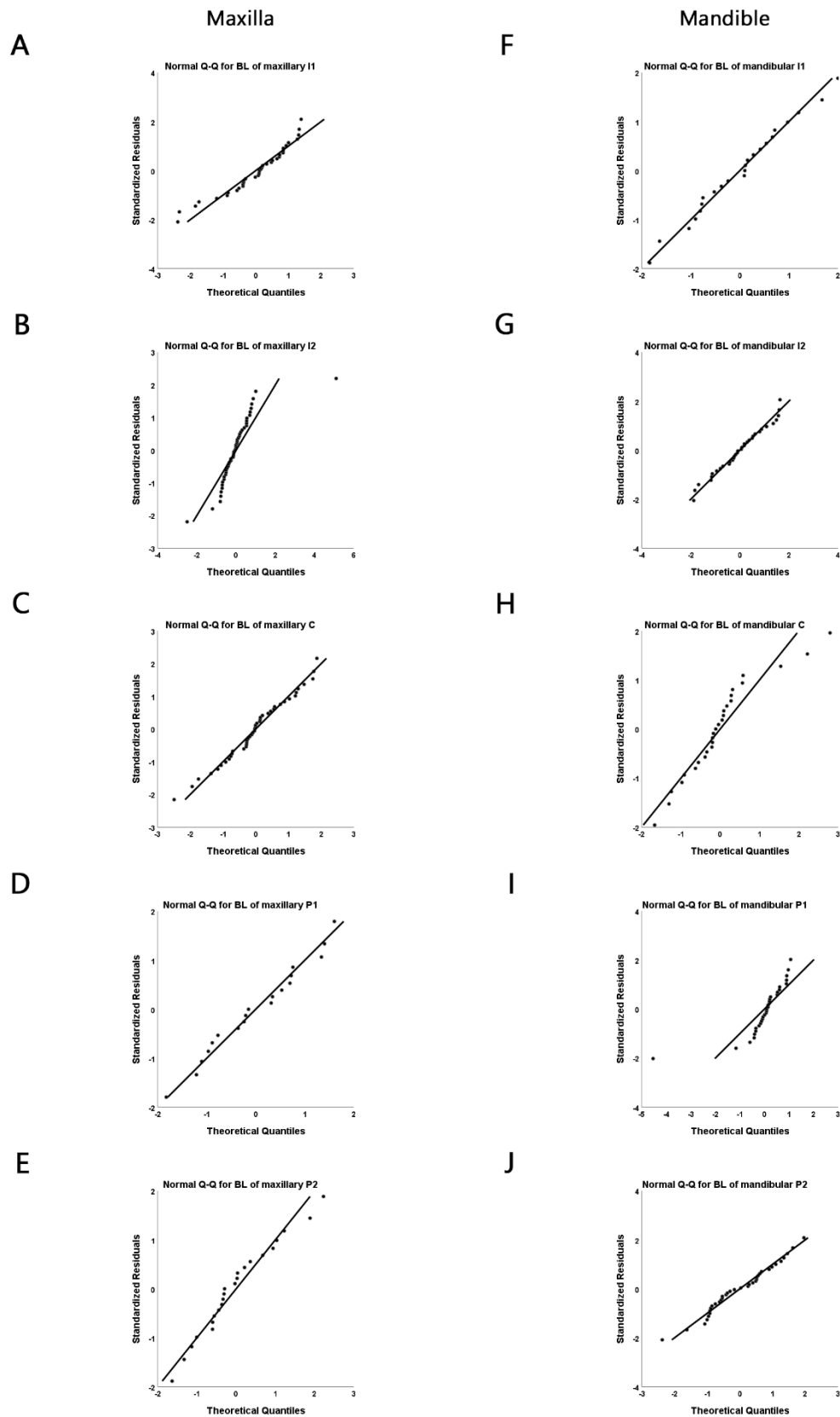


Figure S1e. Q-Q plots (for the models evaluating the **buccolingual cervical diameter**) for the best fit regression models to assess if the set of residuals are normally distributed for the maxillary (A–E) and mandibular (G–J) dentoalveolar structures.

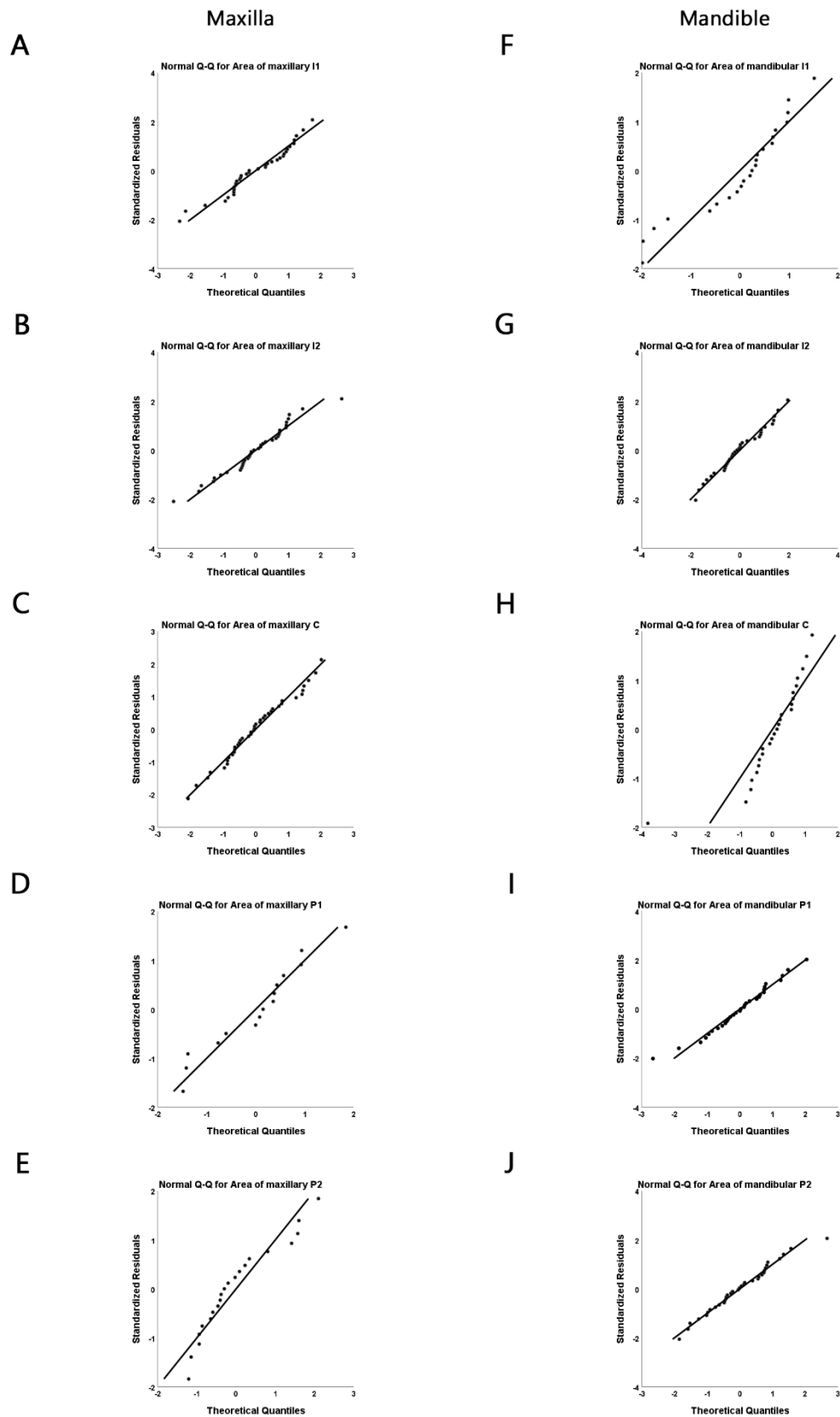


Figure S1f. Q-Q plots (for the models evaluating the **cervical area**) for the best fit regression models to assess if the set of residuals are normally distributed for the maxillary (A–E) and mandibular (G–J) dentoalveolar structures.