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Modeling of muscle forces in humans with and without temporomandibular joint disorders

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Abstract

Objectives—Subjects with/without temporomandibular joint disorders (TMJD) were tested for differences in muscle forces.

Setting and Sample Population—School of Dental Medicine, University at Buffalo. Ninetyone subjects were classified in 4 groups based on presence/absence (+/-) of chronic myofascial and/or TMJ pain (P) and bilateral disc displacement (DD).

Material & Methods—Validated numerical models employed an organizational objective and subjects' anatomy to calculate masticatory muscle forces during static biting. ANOVA and Holm step-down procedure post-hoc tests assessed group differences. Theoretical geometries, representing the range of subjects' muscle orientations, were surveyed via numerical models to identify key combinations resulting in high muscle forces. Effect-size (Cohen's d) and ANOVA/ post-hoc tests assessed group differences in key muscle orientations.

Results—+P-DD subjects had significantly higher muscle forces, especially for lateral pterygoid muscles, compared to the other groups (P<0.01) for bite-forces that were directed posteromedially or posterolaterally on mandibular molars and posteriorly and slightly medially on mandibular incisors. Key muscle orientations for peak lateral pterygoid muscle forces were identified and group comparisons showed mean orientation in +P-DD compared to other diagnostic groups was 5° more upright for masseter and 3° more posteriorly-directed for temporalis muscles (all Cohen's d 0.8).

Conclusion—Predicted lateral pterygoid muscle forces were significantly higher in +P-DD compared to other groups for specific biting conditions and were attributable, in part, to differences in masseter and temporalis muscle orientations.

Keywords

Human; TMJ; masticatory muscles; biomechanics; computer models

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Introduction

Computer-assisted numerical models of human craniomandibular biomechanics (1), using central nervous system (CNS) organizational objectives of minimization of joint loads (MJL) and muscle effort (MME), have proven to be useful tools. For example, these models were employed to elucidate temporomandibular joint (TMJ) eminence growth (2) and shape (3,4), as well as inter-individual differences in TMJ loads (5) and disc mechanics (6). Furthermore, static biting experiments based on data from numerical modeling showed that constants of proportionality between masticatory muscle activities and bite-forces varied with bite-force direction (7,8). These findings suggest simplifying assumptions applied to mathematical modeling such as assignment of muscle forces based on muscle crosssectional areas and averaged electromyographic data from groups of individuals (9,10) are unlikely to yield predictions that can be tested by comparing to individual-specific outcomes. In contrast, results from MJL- and MME-based numerical models (1) have been validated for individuals by comparison with in vivo measurements of TMJ eminence shapes (4) and muscle forces during static biting tasks (11). A next step is to apply these validated numerical models to address clinical questions such as: Do individuals with/without TMJD generate different masticatory muscle forces during biting due to differences in CNS organization and craniomandibular biomechanics? The application of numerical modeling, thus, may translate to predicting groups of individuals who are susceptible to increased jaw muscle activity and joint loading during routine daily function.

Variability amongst diagnostic groups in the biomechanics of biting has been reported (5,6,11). However, it is unknown if, during static biting, ratios of masticatory muscle forces: bite-forces are higher in some clinically defined groups compared to others. Furthermore, it is unknown which anatomical relationships and mandibular loading conditions critically determine biomechanical differences amongst diagnostic groups. Employment of the previously validated numerical models can investigate these unknowns. Outcomes may improve understanding of human susceptibility for development or maintenance of different categories of TMD, and, thus, suggest candidate preventative and treatment approaches.

This project tested the hypothesis that mean predicted masticatory muscle forces during standardized static biting tasks were higher in individuals with (+P), compared to those without, myofascial and/or TMJ pain (-P). Then, using numerical models, anatomical and jaw loading conditions were surveyed to identify those that accounted for the highest masticatory muscle forces. Finally, a second hypothesis was tested to see if these anatomical relations were more common in +P compared to –P individuals.

Material and Methods

Subjects

Institutional Review Boards at University at Buffalo and University of Missouri-Kansas City approved study protocols. The 91 informed, consenting, and qualified subjects (47 women, 44 men) were previously described for the model-validation study (11). Four diagnostic groups were represented according to presence or absence of chronic myofascial and/or TMJ pain (+/-P) and bilateral TMJ disc displacement (+/-DD) determined by a calibrated

examiner and radiologist, respectively using Research Diagnostic Criteria for TMD (12) and computed-tomography and magnetic resonance images (13). Gender was approximately balanced within each group (+P+DD: 13 women, 13 men; +P-DD: 8 women, 8 men; -P +DD: 16 women, 13 men; -P-DD: 10 women, 10 men).

Modeling Protocol and Analyses

Overview of Model Validation—As previously reported (11), individual-specific muscle activation patterns during biting tasks, predicted by computer-assisted numerical models, were tested for accuracy by comparison with masseter and anterior temporalis muscle activities measured in vivo via surface electromyography when the same individual performed similar biting tasks on a bite-force transducer. The transducer was positioned between custom acrylic crowns on maxillary and mandibular right and left central incisors and first molars. The vestubulolingual direction and magnitude of a mechanical moment produced by the bite-force were controlled by orientation of the transducer relative to the center of resistance of each mandibular tooth. For each of four biting positions (left or right, incisors or molars), each subject was asked to produce a range of comfortable bite-forces. For each subject, biting position, moment, and muscle, analyzed data were plotted, slopes were calculated for muscle activity versus bite-force (root mean square mV/N), and normalized to peak slope. Within subjects, normalized results from two laboratory sessions were compared with numerical model-predicted muscle activities during simulation of the in vivo biting tasks. Customized programs calculated differences between measured and predicted data for similar tasks which defined the errors in model results. The modelpredicted and in vivo-measured data with minimum error was identified as the "best-match" result for a given biting task. Average absolute errors between best numerical modelpredicted and in vivo muscle activities were 15% overall and similar amongst diagnostic groups. Errors ranged from 11–13% and 8–15% for incisor and molar biting, respectively. Average coefficients of determination demonstrated that predicted and measured data generally matched well and similarly amongst diagnostic groups, ranging from 0.70–0.74 and 0.68-0.74 for incisor and molar biting, respectively.

Modeling of Muscle Forces during Biting in Diagnostic Groups—Each model employed the subject's anatomical data and an organizational objective to produce unique solutions for static equilibrium. Anatomical data were determined using established methods (4,5,11) and consisted of the subject's three-dimensional craniomandibular geometry (Fig. 1) developed from standardized lateral and postero-anterior cephalometric radiographs and bilateral sagittal effective eminence shapes measured via jaw tracking (4). Objectives were: 1) minimization and equalization of right and left TMJ loads (MJL); or 2) minimization of muscle effort (MME), defined as minimization of the sum of muscle forces squared. Overall, predictions from the MME model fit best with *in vivo* data from incisor and molar biting tasks for 53% of men and women (Table 1). However, for each subject and biting location, the model that predicted muscles forces which matched best with the subject's electromyographic data was used to calculate muscle forces (% applied bite-force) for a spectrum of molar and incisor bite-force angles. Specifically, 324 biting angles were modeled on each of right and left mandibular first molars and central incisors in each subject

according to (Fig. 1): $0-350^{\circ}$ in the occlusal plane (θ_{xz}) in 10° increments, and at each θ_{xz} , $0-40^{\circ}$ relative to vertical (normal to the occlusal plane, θ_{y}) in 5° increments.

Model-predicted forces for each muscle: masseter, temporalis, lateral pterygoid, medial pterygoid, and anterior digastric, relative to the applied bite-force at each biting angle were determined for each subject. Initially, seven variables (gender, diagnostic group, biting location, θ_{xz} , θ_y , muscle, side) were considered. Then, each biting location (molar, incisor) and muscle were considered separately, so omnibus tests of five-factor ANOVAs were conducted using between-subject variables of gender and group (+PP/+DD, +PP/-DD, -PP/+DD, -PP/-DD) and within-subject variables of θ_{xz} (36 angles), θ_y (9 angles) and side (ipsilateral, contralateral to applied bite-force). All interactions up through four-way were examined (five-way was not significant) and where significant, further comparisons with family-wise error testing were used (α =0.05).

Analysis of Geometry Effects on Muscle Forces

Geometries for all subjects were averaged to determine mean positions of mandibular anatomical components important to biting (Fig. 1). Model predictions showed anterior digastric muscles were relatively inactive during biting tasks, as supported by previous in vivo data (8,14) and could, thus be excluded from the analyses. Inspection of the sample population's mean geometry file showed muscle orientations varied most in the sagittal plane (x-y in Fig. 1). Thus, muscle orientations were characterized using mean y-coordinates for each muscle vector, and varying x-coordinates to depict the extremes which ranged from 0 to 45 mm for masseter, -40 to 12 mm for temporalis, 14 to 33 mm for lateral pterygoid, and -4 to 21 mm for medial pterygoid muscles. Each range was represented in the theoretical geometry files by 10 equal steps. A systematic method was applied via customized software (MatLab R2013a, MathWorks, Natick MA), where individual masseter, temporalis, lateral pterygoid, and medial pterygoid muscle orientations were incrementally changed while holding other muscle orientations constant. Through this method, 14,641 theoretical geometries were assembled to reflect the mean geometry with all possible combinations of these ranges of 4 muscle orientations. Theoretical geometry files were used in the two models to identify, through regression analyses, key combinations of muscle orientations that produced the highest predicted muscle forces during biting. These key muscle orientations were then compared amongst groups using effect-size (Cohen's d) and ANOVA with post-hoc tests (α =0.05).

Results

Intergroup Differences in Predicted Muscle Forces during Biting

Polar plots depicted group means in muscle forces for a range of biting angles, where at each θ_{xz} results from θ_y (0–40°) were averaged. Asymmetric shapes of group data sets suggested that bite-force direction, θ_{xz} , was important to muscle forces during biting (Figs. 2 and 3). Generally, +P-DD subjects showed higher predicted forces for all biting conditions tested for ipsilateral and contralateral muscles, as well as masseter, lateral pterygoid and medial pterygoid muscles compared to the other groups (Figs. 2 and 3).

ANOVA showed a substantial number of three-way interactions were significant so paired comparisons with Holm step-down procedures were applied. For posteromedially- or posterolaterally-directed molar bite-forces, predicted muscle forces from individually validated numerical models were significantly higher in the +P-DD compared to the other groups (P<0.01) by: >11% for ipsilateral, >10% for contralateral, and >25% for lateral pterygoid muscles (Fig. 2A–C) and compared to the –P-DD group (P<0.05) by: >15% for masseter (Fig. 2D) and >6% for medial pterygoid (Fig. 2E) muscles. Predicted temporalis muscle forces showed no significant differences between groups for molar biting (data not shown).

Similar comparisons demonstrated that for incisor biting which directed the jaw posteriorly and slightly medially, predicted muscle forces in the +P-DD were significantly higher compared to the other groups (P<0.01) by >26% for ipsilateral muscles (Fig. 3A) and compared to the –P-DD group (P<0.01) by >68% and the +P+DD group (P<0.05) by >55% for lateral pterygoid muscles (Fig. 3B). Predicted contralateral, masseter, medial pterygoid, and temporalis muscle forces showed no significant differences between groups for incisor biting (data not shown).

Survey of Geometry Effects on Muscle Forces during Biting

Geometries which produced the highest predicted lateral pterygoid muscle forces for molar and incisor biting at $\theta_{xz}=30^{\circ}$ and $\theta_{y}=20^{\circ}$ were used to identify key muscle orientations because this muscle and these biting conditions showed the largest between-group differences (Figs. 2 and 3).

The survey showed that peak lateral pterygoid muscle forces for molar and incisor biting predicted by the MJL model were 7.4- and 10.4-fold the applied bite-force, respectively, and by the MME model were 5.6- and 7.9-fold the applied bite-force, respectively (Fig. 4A–D). These peak forces occurred where masseter (x=0mm) and lateral pterygoid (x=14mm) muscles were relatively upright, where temporalis muscles were more posteriorly directed (x=-40, -30mm for MJL, MME models respectively), and were the same for the range of medial pterygoid muscle orientations tested. Step-wise regression analyses confirmed that x-coordinates of masseter, temporalis, and lateral pterygoid muscles accounted for 60% of the variability in predicted lateral pterygoid muscle forces for molar and incisor biting in both models.

Group Differences in Key Muscle Orientations

Key muscle orientations overall were not significantly different (P=0.09) between groups. However, mean masseter muscle orientations were significantly different between +P-DD subjects and the other groups (Table 2; all P 0.05). In +P-DD subjects the masseter muscle was more upright by 5° (Table 3), whereas the temporalis muscle was more posteriorly directed by 3° . Large effect sizes (Table 2, all Cohen's d 0.8) were found for both muscles.

Discussion

+P-DD subjects showed significantly higher predicted muscle forces, especially for lateral pterygoid muscles, compared to other groups for bite-forces directed posteromedially or posterolaterally on mandibular molars and posteriorly and slightly medially on mandibular incisors. These results suggest future tests to see if +P-DD individuals demonstrate higher levels of lateral pterygoid muscle activities and pain compared to others when doing specific biting tasks. *In vivo* data from the lateral pterygoid muscles would also permit an expanded comparison of model-predicted and electromyographic data to validate model results for lateral pterygoid as well as masseter and temporalis muscles. Nevertheless, +P-DD subjects exhibited anatomical differences compared to other groups that could account for differences in jaw biomechanics. Although effect sizes were large, group differences in key muscle orientations overall did not quite reach significant levels (P=0.09), suggesting larger and more balanced sample sizes are indicated in future studies.

Modeling of muscles by single vectors simplifies the complex three-dimensional architecture and localized regional innervation of motor areas. Indeed, some experiments have demonstrated heterogeneity within human masseter (23,24), temporalis (25,26) and lateral pterygoid (27) muscles during jaw loading. However, this occurs with to access to visual feedback during the experiments and/or when teeth with an arch are splinted together. Similar influence of visual feedback on muscle recruitment patterns has been demonstrated in spinal and appendicular muscles (28–30). In contrast, within muscle heterogeneity during jaw loading, does not occur when subjects are without visual feedback and when teeth are not splinted together (31,32). Hence, currently no unequivocal evidence contradicts the use of single vectors to define masticatory muscle orientations for modeling static tasks.

The modeling of muscle forces during static biting was limited to molar and incisor positions. Future work should investigate other jaw loading positions that reflect both functional and non-functional behaviors. Notably, significant group differences occurred at molar biting angles causing distolingual (θ_{xz} =0–60°) and distobuccal (θ_{xz} =290–350°) mandibular crown tipping moments where regional periodontal mechanoreceptor activation provides specific CNS feedback (15). It remains unknown if this range of bite-force directions normally occurs and if certain positions and directions of jaw loading are characteristic for a particular group.

The Pain Adaptation Model (16) suggests that the presence of muscular pain results in inhibition of jaw loading behaviors. However, in a clinical study (11) where subjects with/ without chronic pain performed the same biting tasks, there were no significant differences in magnitudes of voluntary bite-forces, and hence no evidence that pain altered bite-force production. Furthermore, no significant differences were found in high intensity masticatory muscle behavior amongst women with/without myofascial pain during polysomnographic recording (17), whereas low intensity chronic muscle activities were significantly elevated in women with myofascial pain compared to healthy women (18). The more recent Integrated Pain Adaptation Model (19) hypothesizes that individuals will alter neuromuscular organization in particular muscles because of pain. To this end, during *in vivo* biting in baseline compared to experimentally induced pain conditions by otherwise

asymptomatic subjects, patterns of single motor unit firing in the masseter muscle were quickly altered in different ways between different subjects (20). Furthermore, the validated numerical models showed masseter and temporalis muscle activities during biting matched best overall with MME predictions for +/-P and +/-DD groups (11). However, right-left asymmetry in muscle organization was different. For molar biting, asymmetry was common in both TMD (61%) and healthy (53%) subjects, which could reflect habitual sidedness in biting. In contrast, incisor biting showed greater asymmetry in TMD (40%) versus healthy (11%) subjects, which could reflect the phenomenon of altered neuromuscular organization during pain.

From the perspective of a biopsychosocial model of TMD (21,22), biomechanics may be a contributing factor associated with specific TMD diagnostic subgroups. Despite recognized limitations (1), numerical modeling has been validated using *in vivo* data and shown to be a relatively accurate way to study jaw biomechanics in living humans non-invasively (4,11). In addition to the current study, previous work has shown other group differences, such as, in –P+DD subjects, predicted TMJ loads were up to 69% higher compared to healthy subjects (5). Considering this, a next step for application of numerical modeling is the identification of -P-DD individuals with relatively high and low predicted muscle and/or TMJ forces that are linked to high and low TMJ energy densities (6) for long-term follow-up to investigate differences in development of TMD. Further to this, future comparisons of individuals should include additional diagnostic criteria, for example: distinguishing myofascial or TMJ pain or both, TMJ disc position, muscle loading behavior, and heritable traits that affect inflammation and pain sensitivity to increase the likelihood of identifying more homogenous TMD subgroups.

Conclusions

Predicted lateral pterygoid muscle forces were significantly higher in the +P-DD group compared to +P+DD, -P+DD, and –P-DD groups for specific biting conditions and were attributable, in part, to differences in masseter and temporalis muscle orientations.

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Clinical Relevance

Validated computer-assisted numerical models of human jaw mechanics were used to compare muscle forces between people for a comprehensive survey of biting directions on molars and incisors. For the same model, differences in muscle forces during biting between people depend on anatomical differences. Hence, differences in jaw muscle orientation were compared between groups of people with/without TMJDs. Subjects with jaw muscle and/or TMJ pain had anatomical differences that could explain relatively higher muscle forces for the same biting tasks compared to others.



Figure 1.

Axis system and force vectors based on an individual's anatomy used in numerical models (left) including: Applied bite-force (100%, shown at left molar and incisor), joint ($F_{condyle}$), and muscles ($m_{1,2}$ =masseter, $m_{3,4}$ =anterior temporalis, $m_{5,6}$ =lateral pterygoid, $m_{7,8}$ =medial pterygoid, $m_{9,10}$ =anterior digastric muscles). Enlargements (right) show modeled bite-forces characterized by angles in the occlusal plane (θ_{xz} , 0–350°, illustrated above occlusal plane for left biting angles; right biting angles were mirror images) and relative to vertical (θ_y , 0–40° where 0° is normal to the occlusal plane). Modified from (11).



Figure 2.

Mean predicted muscle forces during molar biting in 4 diagnostic groups with/without (+/-) pain (P) and TMJ disc displacement (DD). Circumferential scale indicates biting angle in occlusal plane (θ_{xz} , Fig. 1). Radial scales indicate muscle force magnitude (%) relative to applied bite-force (100%). The +P-DD group showed significantly higher: (A) Ipsilateral muscle forces than other groups (all P<0.01, red area): θ_{xz} =0–100° and 270–350°; (B) Contralateral muscle forces than other groups (all P<0.01, red area): θ_{xz} =0–60° and 290–350°; (C) Lateral pterygoid muscle forces than other groups (all P<0.01, red area): θ_{xz} =0–60° and 290–90° and 260–350°; (D) Masseter muscle forces than the –P-DD group (P<0.05, green area): θ_{xz} =0–90° and 290–350°; and (E) Medial pterygoid muscle than the –P-DD group (P<0.05, green area): θ_{xz} =0–90° and 290–350;



Figure 3.

Mean predicted muscle forces during incisor biting in 4 diagnostic groups with/without (+/-) pain (P) and TMJ disc displacement (DD). Circumferential scale indicates biting angle in occlusal plane (θ_{xz} , Fig. 1). Radial scales indicate muscle force magnitude (%) relative to applied bite-force (100%). The +P-DD group showed significantly higher (A) Ipsilateral muscle forces than other groups (all P<0.01, red area) for: θ_{xz} =10–40°; and (B) Lateral pterygoid muscle forces than the –P-DD (P<0.01, red area) and +P+DD (P<0.05, stippled area) groups for: θ_{xz} =0–40° and 350°.



Figure 4.

Theoretical analysis of muscle orientation effects on lateral pterygoid muscle forces. 3D plots of MJL (A,C) and MME (B,D) predicted lateral pterygoid muscle forces for molar (A,B) and incisor (C,D) biting, each with a unique color scale. Orientation ranges for each muscle found in the sample were used in two models to identify key combinations of muscle orientations resulting in the highest predicted muscle forces during biting. Lateral pterygoid muscle forces are expressed relative to a 100 unit bite-force (%). Step-wise regression analyses confirmed that x-coordinates of masseter, temporalis, and lateral pterygoid muscles accounted for 60% of predicted lateral pterygoid muscle forces increased when: i) masseter muscles were more upright; ii) temporalis muscles were more posteriorly directed; and iii) lateral pterygoid muscles were less anteriorly directed.

Table 1

Frequency distribution of "best model" for predicting muscle activities during molar and incisor biting in men and women. The data from all groups are combined, where MME (minimization of muscle effort) or MJL (minimization of joint load) were the objective functions used by the computer models to predict *in vivo* muscle activities. Model predicted data were compared with *in vivo* data. On average, model accuracy was 86–91% for predicting these *in vivo* muscle activities (11).

Gender	Biting Position	MME (%)	MJL (%)
Men	Molar	81	19
Women	Molar	72	28
Men	Incisor	72	28
Women	Incisor	53	47

Table 2

Comparisons of mean muscle x-coordinates for +P-DD group with other groups. Cohen's d effect size and ANOVA/post-hoc tests were used to characterize differences.

Muscle	Cohen's d effect size (P-value, unadjusted post-hoc test):				
	+P-DD vs +P+DD	+PP-DD vs -P+DD	+PP-DD vs -P-DD		
Masseter	2.2 (0.02)	2.0 (0.02)	2.2 (0.05)		
Temporalis	1.1 (0.23)	0.8 (0.32)	1.6 (0.07)		
Lateral Pterygoid	1.4 (0.09)	0.4 (0.72)	0.4 (0.69)		
Medial Pterygoid	0.5 (0.64)	0.2 (0.82)	0.3 (0.74)		

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Table 3

Mean (standard deviation) of x-coordinates (mm) for muscles in diagnostic groups.

Diagnostic Group	Masseter	Temporalis	Lateral Ptyergoid	Medial Pterygoid
+P+DD (n=26)	27.8 (6.8)	-0.1 (5.6)	24.2 (4.0)	8.5 (5.2)
+P-DD (n=16)	22.3 (9.0)	-3.4 (12.8)	22.4 (4.0)	9.3 (5.2)
-P+DD (n=29)	27.8 (8.9)	-0.7 (9.3)	22.0 (2.5)	9.7 (5.8)
-P-DD (n=20)	27.4 (4.5)	2.0 (6.2)	22.8 (2.7)	8.7 (5.4)