

## Fixed Apparatus Medication of a Skeletal Class II Open Chomp Malocclusion with Hypomature Amelogenesis Blemished: Case Report

Halimi A<sup>1\*</sup>, El alloussi M<sup>2</sup>, Sefiani A<sup>3</sup> and Zaoui F<sup>1</sup>

<sup>1</sup>University Mohammed V-Souissi (UM5S), Faculty of Dentistry, Biotech Research Team. MMB, Ibn Sina Hospital, Centre for Dental Consultation and Treatment, Department of Orthodontics and dentofacial Orthopedics, Morocco

<sup>2</sup>University Mohammed V-Souissi (UM5S), Faculty of Dentistry Rabat Ibn Sina Hospital, Centre for Dental Consultation and Treatment, Department of Pediatric dentistry, Morocco

<sup>3</sup>University Mohammed V-Souissi (UM5S), genomic center of human science, Department of medical genetic, Morocco

### Abstract

This article depicts the orthodontic medicine of a 12-year-old female tolerant having a skeletal Class II malocclusion with localized hypomature amelogenesis imperfecta. We were fit to fulfil orthodontic medication without causing dental polish harm for example demineralization, white spot sore and breaks when were moved the edgewise apparatuses. A great impediment and feel grins were realized, and these comes about have been administered for three years after fruition of the animated medication.

**Keywords:** Altered apparatus medicine; Class II open chomp malocclusion; Hypomature amelogenesis imperfecta

### Introduction

In an instance of extreme skeletal Class II malocclusion with long challenge, we frequently have trouble regulating the posterior movement of the molars throughout orthopaedic medication. Since the patient had an extraordinary mellow localized manifestation of hypomature amelogenesis imperfecta, we established to build sufficient plaque control and treated the patient by a straight holding strategy.

Amelogenesis imperfecta (AI) has been portrayed as a mind boggling assembly of inherited conditions that exasperates the improving polish structure and exists free of any identified systemic disorder [1-4]. This lacquer abnormality influences both the essential and changeless dentition [1-5]. The occurrence of amelogenesis imperfecta has been accounted for to differ between give or take 1:700 and 1:16,000, relying on the populace considered and the analytic criteria used [5-8].

The most acknowledged characterization framework recognizes three fundamental AI sorts dependent upon the anticipated developmental system: hypoplastic (HPAI)-secretory deformity; hypocalcified (HCAI)-crystallite nucleation and development imperfection; and hypomaturation (HMAI)-protein transforming and crystallite development deformity. Further divisions are dependent upon mode of inheritance [9,10].

HPAI comes about because of deformities in the secretory technique making slender or hollowed veneer that could be typical or adjusted in structure or alternately creation. Hypoplastic AI is connected with various allelic transformations in the AMELX (OMIM 606585) and ENAM (OMIM 300391) genes and presumable will indicate heterogeneity past the aforementioned genes [11-18].

The aetiology of HCAI is accepted to be a powerlessness of the crystallites to fittingly nucleate initiating anomalous and a stamped diminishing in mineral substance. The sub-atomic surrender for HCAI remains obscure. Nonetheless, in a few expansive families, the most ordinarily recognized AI petitioner genes have been prohibited through linkage investigation showing up 'til now unidentified genes are connected with this AI type [19].

HMAI is brought about by strange of the lattice proteins throughout development coming about because of either irregular of veneer mutant framework proteins (e.g. AMELX changes) or anomalous (e.g. KLK4 transformations) [12,20,21]. The expanded protein maintenance in HMAI avoids the ordinary advancement and development of the finish gems and brings about a diminished polish mineral substance. Change in the kallikrein 4 gene (KLK4) that transforms a finish serine proteinase basic for veneer development has been distinguished in one family having autosomal passive pigmented HMAI [21].

A Swedish investigation of 66 individuals with AI discovered skeletal open nibble qualities in families with X-connected AI, autosomal latent summed up flimsy HPAI and AI subtypes described by hypomineralization (i.e. HCAI and HMAI) [22].

The pervasiveness of open nibble malocclusions in individuals with lacquer deserts diagnosed as AI is more excellent than in the all inclusive community. In any case, the commonness of open nibble in AI relatives without polish absconds is additionally more stupendous than the general population [22]. While it is plausible that the hereditary changes answerable for AI finish imperfections either incline a single person to or creates an open nibble, it moreover is conceivable that the AI cohorted veneer absconds and open nibbles are brought on by distinctive hereditary systems.

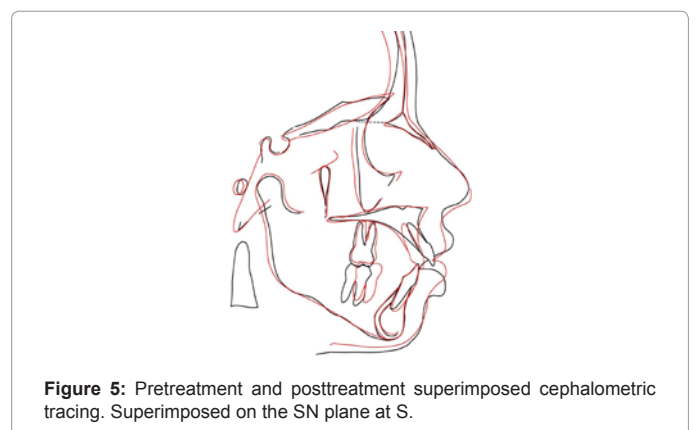
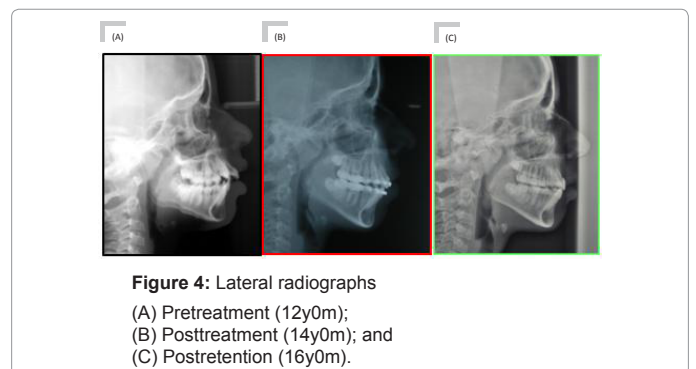
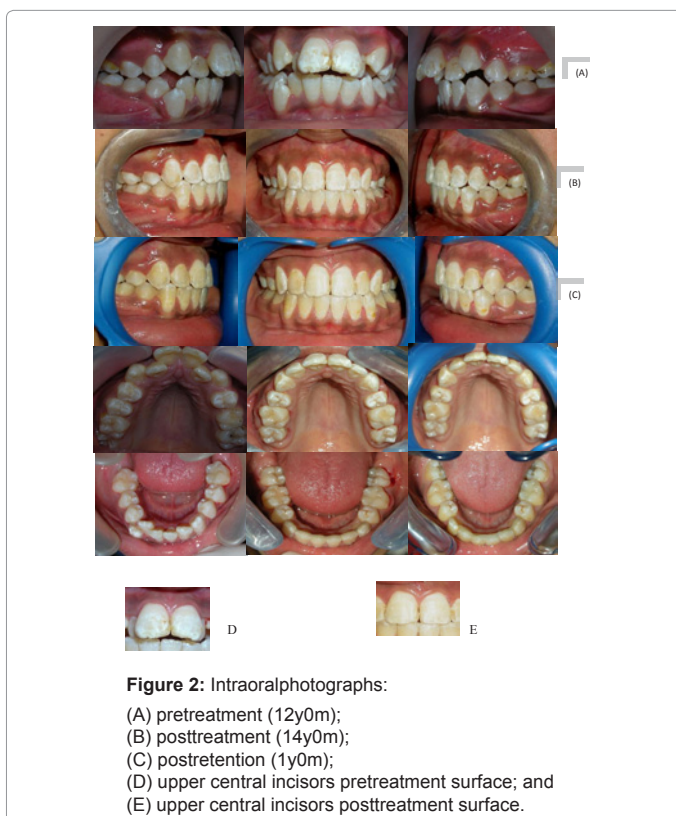
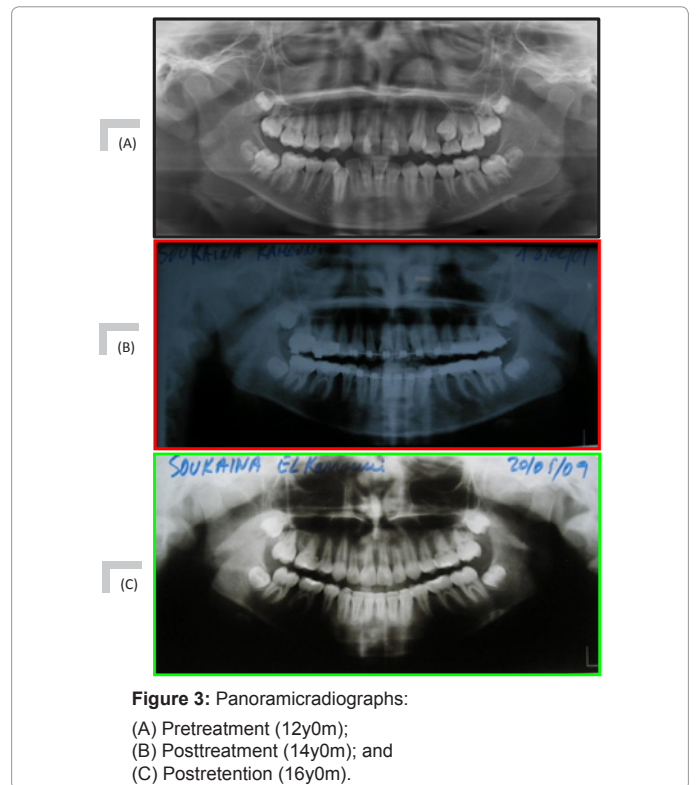
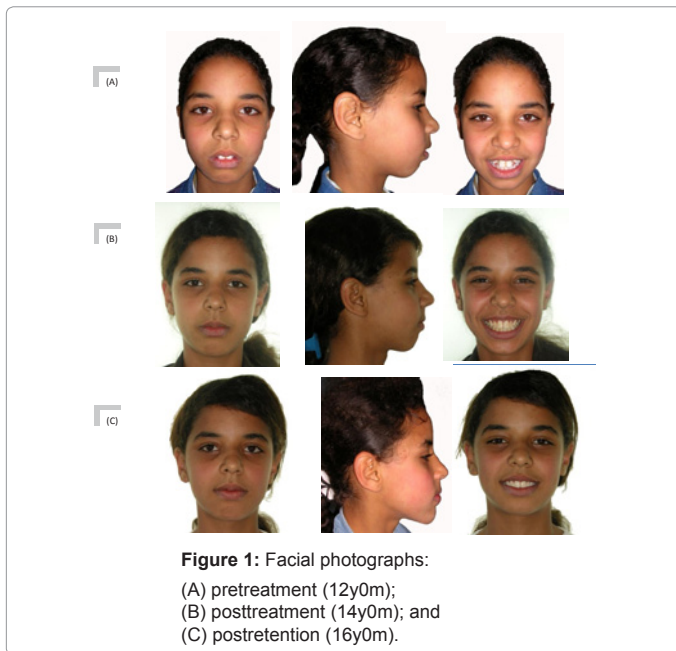
Cartwright et al. [23] Assessed open chomp attributes in AI relatives that did not express veneer deserts as controls to separate

**\*Corresponding author:** Dr. Abdelali Halimi, Orthodontist, MD, PhD, University Mohammed V Souissi, Faculty of Dentistry Rabat, Morocco, Tel: +212666285639, 00 212 37790844; E-mail: [halimiali111@yahoo.fr](mailto:halimiali111@yahoo.fr)

Received April 09, 2013; Accepted May 03, 2013; Published May 05, 2013

**Citation:** Halimi A, El alloussi M, Sefiani A, Zaoui F (2013) Fixed Apparatus Medication of a Skeletal Class II Open Chomp Malocclusion with Hypomature Amelogenesis Blemished: Case Report. J Nov Physiother 3: 137. doi:10.4172/2165-7025.1000137

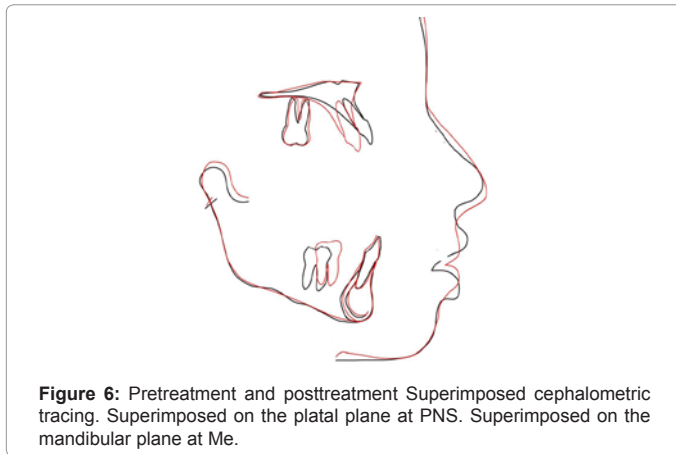
**Copyright:** © 2013 Halimi A, et al. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.



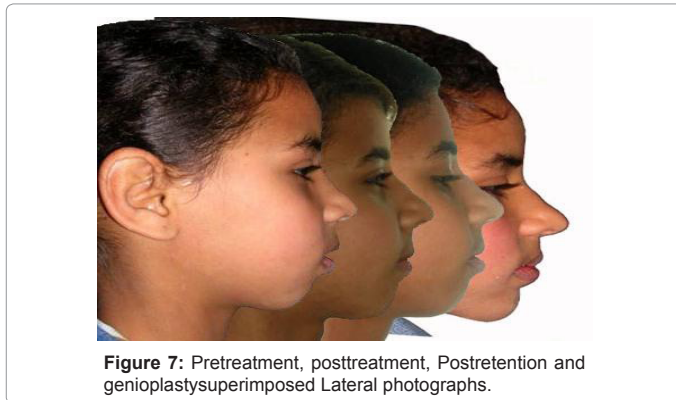
if the open nibble phenotype was a familial quality free of AI [22]. In some AI kindred's, the skeletal open nibble happened in relatives with and without veneer deserts proposing that, in anyhow some AI families, the aforementioned two phenotypic attributes could be disconnected. All things considered, the aforementioned clinical studies propose an in number yet as yet poorly demarcated acquaintanceship between the presence of AI veneer imperfections and open nibble malocclusions. Understanding this companionship

is clinically significant as numerous intense AI open chomps need surgical-orthodontic correction [23,24].

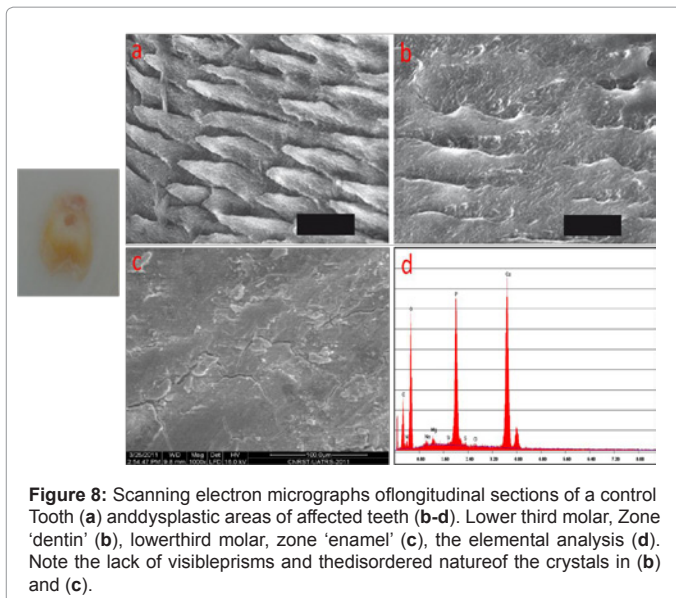
The pathophysiology prompting an affiliation between dental or



**Figure 6:** Pretreatment and posttreatment Superimposed cephalometric tracing. Superimposed on the palatal plane at PNS. Superimposed on the mandibular plane at Me.



**Figure 7:** Pretreatment, posttreatment, Postretention and genioplastysuperimposed Lateral photographs.



**Figure 8:** Scanning electron micrographs of longitudinal sections of a control Tooth (a) and dysplastic areas of affected teeth (b-d). Lower third molar, Zone 'dentin' (b), lower third molar, zone 'enamel' (c), the elemental analysis (d). Note the lack of visible prisms and the disordered nature of the crystals in (b) and (c).

alternately skeletal open nibble malocclusion and AI remains unclear [22,25-28].

## Case Presentation

The patient was a 10-year-old Marocain female with an intense skeletal Class II open nibble malocclusion, tongue brokenness. The head dissention was the maxillary bulge and gathering. In her family

history, her father had amelogenesis imperfecta.

The patient had a raised profile, with abundance vertical stature of the easier front side (Figure 1). Intraorally she had an Angle Class II molar of 7 mm relationship in the right side, Angle Class II molar of 6 mm relationship in the left agree with, a minimal overbite of-1 mm (between 14 and 24) and an over jet of +7 mm. The premolars and molars were positioned in crossbite. The bend of Spee was-3 mm and gathering-6 mm was available in the mandibular curve. The mandibular midline moved to the left by 2 mm to the facial one. Dental curve width of maxillary and mandibular showed disharmony. The hypomature amelogenesis imperfecta localised at the upper incisors (Figure 2).

The surrounding radiographs demonstrated blended dentition, presence of the 65 and germs of all third molars (Figures 3- 8).

Cephalometric dissection indicated a skeletal Class II association with an ANB plot of 8°. The measure of the mandible was minor and it was ordinary.

## References

- Weinmann JP, Svoboda JF, Woods RW (1945) Hereditary disturbances of enamel formation and calcification. J Am Dent Assoc 32: 397-418.
- Aldred MJ, Savarirayan R, Crawford PJ (2003) Amelogenesis imperfecta: a classification and catalogue for the 21st century. Oral Dis 9: 19-23.
- Neville BW, Damm DD, Allen CM, Bouquot JE (2002) Oral and maxillofacial pathology 2nd ed. Philadelphia: Elsevier 89-94.
- Robinson FG, Haubenreich JE (2006) Oral rehabilitation of a young adult with hypoplastic amelogenesis imperfecta: a clinical report. J Prosthet Dent 95: 10-13.
- Rao S, Witkop CJ Jr (1971) Inherited defects in tooth structure. Birth Defects Orig Artic Ser 7: 153-184.
- Bäckman B, Holm AK (1986) Amelogenesis imperfecta: prevalence and incidence in a northern Swedish county. Community Dent Oral Epidemiol 14: 43-47.
- Sundell S, Koch G (1985) Hereditary amelogenesis imperfecta. I. Epidemiology and clinical classification in a Swedish child population. Swed Dent J 9: 157-169.
- WITKOP CJ (1957) Hereditary defects in enamel and dentin. Acta Genet Stat Med 7: 236-239.
- Ravassipour DB, Powell CM, Phillips CL, Hart PS, Hart TC, et al. (2005) Variation in dental and skeletal open bite malocclusion in humans with amelogenesis imperfecta. Arch Oral Biol 50: 611-623.
- WITKOP CJ (1957) Hereditary defects in enamel and dentin. Acta Genet Stat Med 7: 236-239.
- Witkop CJ Jr (1988) Amelogenesis imperfecta, dentinogenesis imperfecta and dentin dysplasia revisited: problems in classification. J Oral Pathol 17: 547-553.
- Wright JT, Hart PS, Aldred MJ, Seow K, Crawford PJ, et al. (2003) Relationship of phenotype and genotype in X-linked amelogenesis imperfecta. Connect Tissue Res 44 Suppl 1: 72-78.
- Hart PS, Michalec MD, Seow WK, Hart TC, Wright JT (2003) Identification of the enamelin (g.8344delG) mutation in a new kindred and presentation of a standardized ENAM nomenclature. Arch Oral Biol 48: 589-596.
- Lagerström M, Dahl N, Nakahori Y, Nakagome Y, Bäckman B, et al. (1991) A deletion in the amelogenin gene (AMG) causes X-linked amelogenesis imperfecta (AIH1). Genomics 10: 971-975.
- Aldred MJ, Crawford PJ, Roberts E, Thomas NS (1992) Identification of a nonsense mutation in the amelogenin gene (AMELX) in a family with X-linked amelogenesis imperfecta (AIH1). Hum Genet 90: 413-416.
- Lench NJ, Winter GB (1995) Characterisation of molecular defects in X-linked amelogenesis imperfecta (AIH1). Hum Mutat 5: 251-259.
- Mårdh CK, Bäckman B, Holmgren G, Hu JC, Simmer JP, et al. (2002) A

- nonsense mutation in the enamelin gene causes local hypoplastic autosomal dominant amelogenesis imperfecta (AIH2). *Hum Mol Genet* 11: 1069-1074.
18. Rajpar MH, Harley K, Laing C, Davies RM, Dixon MJ (2001) Mutation of the gene encoding the enamel-specific protein, enamelin, causes autosomal-dominant amelogenesis imperfecta. *Hum Mol Genet* 10: 1673-1677.
19. Hart PS, Wright JT, Savage M, Kang G, Bensen JT, et al. (2003) Exclusion of candidate genes in two families with autosomal dominant hypocalcified amelogenesis imperfecta. *Eur J Oral Sci* 111: 326-331.
20. Ravassipour DB, Hart PS, Hart TC, Ritter AV, Yamauchi M, et al. (2000) Unique enamel phenotype associated with amelogenin gene (AMELX) codon 41 point mutation. *J Dent Res* 79: 1476-1481.
21. Hart PS, Hart TC, Michalec MD, Ryu OH, Simmons D, et al. (2004) Mutation in kallikrein 4 causes autosomal recessive hypomaturational amelogenesis imperfecta. *J Med Genet* 41: 545-549.
22. Bäckman B, Adolfsson U (1994) Craniofacial structure related to inheritance pattern in amelogenesis imperfecta. *Am J Orthod Dentofacial Orthop* 105: 575-582.
23. Cartwright AR, Kula K, Wright TJ (1999) Craniofacial features associated with amelogenesis imperfecta. *J Craniofac Genet Dev Biol* 19: 148-156.
24. Wright JT, Waite P, Mueninghoff L, Sarver DM (1991) The multidisciplinary approach managing enamel defects. *J Am Dent Assoc* 122: 62-65.
25. Hoppenreijts TJ, Voorsmit RA, Freihofer HP (1998) Open bite deformity in amelogenesis imperfecta. Part 1: An analysis of contributory factors and implications for treatment. *J Craniomaxillofac Surg* 26: 260-266.
26. Witkop CJ, Sauk JJ (1976) Heritable Defects of Enamel. In: Stewart R, Prescott G, editors. *Oral facial genetics*. St. Louis: C.V. Mosby Company 151-226.
27. Rowley R, Hill FJ, Winter GB (1982) An investigation of the association between anterior open-bite and amelogenesis imperfecta. *Am J Orthod* 81: 229-235.
28. Nishimura K, Hidaka K, Kitagawa H, Goto S (2006) Orthodontic correction of a skeletal Class III malocclusion with impacted maxillary second molars and amelogenesis imperfecta orthodontic waves 65: 43-49.

**Citation:** Halimi A, El alloussi M, Sefiani A, Zaoui F (2013) Fixed Apparatus Medication of a Skeletal Class II Open Chomp Malocclusion with Hypomature Amelogenesis Blemished: Case Report. J Nov Physiother 3: 137. doi:[10.4172/2165-7025.1000137](https://doi.org/10.4172/2165-7025.1000137)

### Submit your next manuscript and get advantages of OMICS Group submissions

#### Unique features:

- User friendly/feasible website-translation of your paper to 50 world's leading languages
- Audio Version of published paper
- Digital articles to share and explore

#### Special features:

- 250 Open Access Journals
- 20,000 editorial team
- 21 days rapid review process
- Quality and quick editorial, review and publication processing
- Indexing at PubMed (partial), Scopus, EBSCO, Index Copernicus and Google Scholar etc
- Sharing Option: Social Networking Enabled
- Authors, Reviewers and Editors rewarded with online Scientific Credits
- Better discount for your subsequent articles

Submit your manuscript at: <http://www.omicsonline.org/submission/>

