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Relationships Between Cranial Base Synchondroses and Craniofacial Development: A Review

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Abstract: Synchondrosis is defined as the development of a union between two bones by the formation of either hyaline cartilage or fibro-cartilage. This paper reviews the relationship between cranial base synchondroses and craniofacial development. The cranial base synchondroses are important growth centers of the craniofacial skeleton. Their abnormalities lead to numerous growth and developmental conditions in the craniofacial region. In dentofacial orthopedics, mechanical forces are commonly applied to cranial bones for growth modification to treat such conditions. Molecular biology and genetics provide tools to investigate the molecular mechanisms, genes and transcription factors responsible for synchondrosis and craniofacial development.

Keywords: Synchondrosis, craniofacial growth.

INTRODUCTION

Cranial base synchondroses are important growth centers of the craniofacial skeleton and the last sites in the cranium to terminate growth [1-5]. Abnormalities in synchondroses are the cause of numerous growth and developmental conditions of the craniofacial region, such as Apert syndrome (1:160000 to 1:200000 live births) [6], Crouzon syndromes (1:60000 live births) [7], Pfeiffer syndromes (1:100000 live births) [8], and Down syndrome (13:10000 live births) [9]. Some abnormalities are craniosynostoses, or premature bony fusions of skull sutures, which eventually influence the life quality of affected children and their families [10]. Much effort has been made to overcome these problems. For example, in dentofacial orthopedics, mechanical forces are commonly applied to cranial bones for growth modification [11-14].

A search of Medline Ovid from 1950 identified 258 papers on synchondrosis, consisting of 188 papers written in English and 70 in other languages, such as German, French, Japanese, Italian, Polish, Czech and Russian. Most of the papers were case reports and some were experimental reports, but only a few were literature reviews. In the light of this finding, this paper reviews the literature on cranial base synchondroses, particularly the sphen-ethmoidal synchondrosis, focusing on sources written in English either as e-journals or in printed journals or textbooks, which were hand-searched.

DEFINITION

Synchondrosis is defined as the development of a union between two bones by the formation of either hyaline cartilage or fibro-cartilage. A synchondrosis is usually temporary and exists during the growing phase until the intervening cartilage becomes progressively thinner during skeletal maturation and ultimately becomes obliterated and converted into bone before adult life. In simple terms, a synchondrosis is a cartilaginous joint. It allows only slight movement between bones compared with the synovial joint, which has a much greater range of movement. Throughout the human body, cartilaginous joints can be found in the pubic symphysis, between the ribs and sternum, between adjacent vertebrae in the spinal column, and at the growth plates between the epiphysis and metaphysis of tubular bones. Three synchondroses are present along the midline of the cranial base: the sphen-ethmoidal synchondrosis between the sphenoid and ethmoidal bones, the intersphenoid synchondrosis between two parts of the sphenoid bone and the sphen-occipital synchondrosis between the sphenoid and basioccipital bones.

Cranial base synchondroses are regarded as important growth centers of the craniofacial skeleton, particularly the sphen-occipital synchondrosis because of its late ossification and important contribution to post-natal cranial base growth [1, 15-18]. The sphen-ethmoidal synchondrosis terminates at 6 years of age [15, 19] and the intersphenoid synchondrosis ossifies immediately before birth [20]. The sphen-occipital synchondrosis seems to have a more prominent role in growth (ontogeny) of the human skull up to adult life than other midline chondral structures [19]. Synchondroseal cartilage is abundant at birth and contains a zone of proliferative and hypertrophic cells that show active growth up to 3 to 4 years of age, after which reduction in growth is observed [21].

HISTOLOGICAL STRUCTURE AND GROWTH MECHANISM OF SYNCHONDROSES

Morphologically, a synchondrosis is similar to the long-bone growth plate, except that growth at the synchondrosis is not unipolar but bipolar. Indeed, the synchondrosis can be regarded as two growth plates positioned back to back so that they share a common zone of actively proliferating
chondroblasts, or the “rest zone”. The different zones of the
synchondrosis mirror each other such that there is cartilage
in the center and bone at each end. The intraoccipital
synchondrosis is completely ossified by 5 weeks of postnatal
development, whereas the intrasphenoid and sphen-
occipital synchondrosis retain a small amount of cartilage
into adulthood.

Specific characteristics of several functional cell zones of
a synchondrosis can be easily identified (Fig. 1) [22]:

1. Zone of vascular erosion and invasion, 1 to 2 cells
   thick
2. Zone of chondrolacunar hypertrophy and matrix
calcification, 2 to 3 cells thick
3. Zone of matrix production or matrixogenic zone, 4 to
   6 cells thick
4. Zone of cellular proliferation, 4 to 6 cells thick
5. Central zone or resting zone, 10 to 12 cells thick

The distinct columnar organization of zones 2 to 4 is
analogous to the organization of the long-bone epiphyseal
plates, which are characterized structurally by bipolar
epiphyseal cartilage. These zones are responsible for the
growth mechanism of endochondral bone, with the central
zone as the true synchondrosis. The central, or resting, zone
is synonymous to the reserve zone of the growth plate and is
composed of chondrocyte precursors that direct the
formation and organization of a synchondrosis. The cells in
the central zone are much denser and smaller than cells in
any of the other layers; they are also not arranged in
columns, and are surrounded by a thicker layer of cartilage
matrix [23].

The proliferative zone plays a crucial role in
endochondral bone formation and is the region of active cell
replication [24]. In the proliferating layer, the chondrocytes
are arranged in columns that are parallel to the longitudinal
axis of the bone and are separated from each other by
matrices containing large amounts of collagen type II. The
number of cells in each column ranges from 10 to 20,
depending on the mitotic moment of the column. The more
cells there are, the greater the growth rate is [25].
Furthermore, multiple mitoses can be observed in the
proximal portion of each column, even though each column
behaves independently [26]. Because the two daughter cells
line up along the long axis of the bone when the
chondrocytes divide, clones of chondrocytes are arranged in
columns parallel to this axis. This spatial orientation directs
growth in a specific direction and is thus responsible for the
elongate shape assumed by many endochondral bones. The
chondrocytes divide only in this zone of the growth plate,
and the high degree of division makes the cells flat and
slightly irregular in shape. The proliferating layer has a high
oxygen content, and the chondrocytes store large amounts of
glycogen and have a high rate of mitochondrial ATP
production [27]. The matrix is synthesized in the lower half
of the proliferating layer by an extensive network of

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Fig. (1). Photomicrograph of a synchondrosis in neonatal mice showing different zones across one side of the synchondrosis. (A) Zone of vascular erosion and invasion. (B) Zone of chondrolacunar hypertrophy and matrix calcification. (C) Zone of matrix production or matrixogenic zone. (D) Zone of cellular proliferation. (E) Central zone or resting zone.
chondrocytes. Ultrastructural examination has shown that these chondrocytes have large amounts of rough endoplasmic reticulum, while an autoradiographic study using tritium-labeled thymidine has revealed high rates of cell division in the three to six layers of flattened cells of the proliferative zone, with lower rates in both the resting/central zone and the prehypertrophic or matrixogenic zone [28, 29].

The hypertrophic zone also plays a key role in endochondral bone formation. Hypertrophic chondrocytes are generated by terminal differentiation of the proliferative zone chondrocytes farthest from the epiphysis. The function of the hypertrophic cell layer is not only to synthesize matrix but also to cause growth by cell expansion/hypertrophy [30-32]. The hypertrophic chondrocyte, by the mechanism of swelling, is actually the principal engine of longitudinal bone growth. During the differentiation process, the chondrocytes increase their intracellular volume by 5 to 10 times, a sign of major activity in the intracellular organelles and cytosol. These cells cease dividing and then enlarge, contributing substantially to the growth process [33]. The hypertrophic chondrocyte is a viable master regulatory cell with an active metabolism. It initiates ossification, which prepares the matrix for calcification and vascularization by attracting vascular and bone-cell invasion from the adjacent bone [34].

In contrast to the proliferative and hypertrophic zones, the resting zone has an unknown function. Abad and coworkers [35] suggested that resting zone cartilage makes important contributions to endochondral bone formation at the growth plate by providing stem-like cells to produce proliferative chondrocytes and producing growth plate-orienting factors (GPOF), which are morphogens that control the alignment of the proliferative clones into columns parallel to the long axis of the bone. The resting zone also probably produces another type of morphogen that is responsible for inhibiting terminal differentiation of the nearby proliferative zone and thus the organization of the growth plate into distinct zones of proliferation and hypertrophy.

**EVOLUTIONARY SIGNIFICANCE OF CRANIAL BASE IN CRANIOFACIAL DEVELOPMENT**

The cranial base has been an interesting subject of research owing to its importance in integrated craniofacial development and growth. It is the most complex structure of the human skeleton and its main function is to protect and support the brain. The cranial base is also the first region of the skull to reach adult size and is the structural foundation of many aspects of the craniofacial architecture [36]. In addition, it also connects with the rest of the body, articulates with the vertebral column and the mandible, provides conduits for all the vital neural and circulatory connections between the brain and the face and neck, houses and connects the sense organs in the skull and forms the roof of the nasopharynx. The shape of the cranial base is therefore a multifactorial product of numerous phylogenetic, developmental and functional interactions [37, 38].

Although the cranial base, vault and face derive from embryologically distinct regions (basocranium, neurocranium and splanchnocranium), the cranial base region grows in a morphologically integrated manner through numerous developmental and functional interactions [39-41]. These interactions occur as a result of many morphogenetic (e.g., neural) and functional (e.g., masticatory, respiratory) stimuli, but the role of the cranial base in influencing overall cranial shape merits special consideration. The cranial base is derived from the chondrocranium, which serves as a growth plate, and is later replaced by bone through endochondral ossification. The chondrocranium is essentially a structural template within and around which the bony cranial base is constructed [42]. Any cartilage remaining between the bones in the form of synchondroses acts as a growth center in a similar way to the epiphysial plates of the long bones. By the end of adolescent growth, the cranial base synchondroses have completely ossified and longitudinal growth has finished. Any interference that causes early or late ossification will result in shortening or lengthening of the cranial base, respectively, leading to serious consequences in craniofacial development [22].

**BIOLOGICAL BASIS OF CRANIAL BASE SYNCHONDROSSES**

Information on the cartilaginous joint in the cranial base is limited, probably as a result of general acceptance that basal synchondroses are the same as the epiphysial plates of the long bones [43]. Both share the same internal morphological features, as well as roles in endochondral bone growth [44]. In the postnatal period, endochondral ossification of the cranial base synchondroses contributes to the expansion of the ossification centers and growth of the cranial base. Therefore, cranial base synchondroses are analogous structures to growth plates of the long bones, in that they allow for the rapid endochondral growth of the bone and eventually close when the bone reaches its final size. Typical long-bone growth cartilage has been defined as a simple proliferating system that exhibits a linear mathematical relationship between cell division and bone growth [45]. Although the overall development of cranial base synchondroses has received less attention than that of long-bone growth plates, many genes have been identified that have proven to be fundamental in the development of the craniofacial complex [46].

**EMBRYOLOGICAL DEVELOPMENT OF CRANIAL BASE SYNCHONDROSSES**

The cranial base has a distinct pattern of morphogenesis. The anterior cranial base is derived from neural crest mesoderm, which is also responsible for the growth of facial bone. In contrast, the posterior cranial base is developed from paraxial mesoderm. The cranial base develops from a cartilaginous template that is replaced by bone through the process of endochondral ossification. The cranial base is initially condensed, as individual elements. Pairs of templates, from caudal to rostral regions, are composed of scleroticome cartilage, parachordal cartilage (precursor of the basioccipital bone), hypophyseal cartilage (precursor of the basisphenoid), presphenoid cartilage (precursor to the sphenoid body anterior to the tubeculum sellae and chiasmatic sulcus), orbitosphenoid (precursor of anterior clinoid and lesser wing of the sphenoid), alisphenoid cartilage (precursor of greater wing of sphenoid) and mesethmoid cartilage. Ossification progresses in an orderly pattern from the posterior to the anterior and finally fuses into a single, irregular and much-perforated base plate,
known as the chondrocranium [47]. The chondrocranium then undergoes further progressive ossification. Large areas of the cranial base remain cartilaginous throughout early fetal life, while many cartilages persist into the third trimester and postnatal life in humans [10].

Histological studies indicate that at least the sphenopalatine synchondrosis contains cartilaginous canals through which blood vessels travel, in addition to connective tissue elements, probably for the purpose of bringing nutrients and other supplies to the cell [48]. Several in vivo studies have been designed to investigate the development of cranial base synchondroses in using mice fetuses [49-51]. The basioccipital synchondrosis was the first to form, starting at 14 days post coitum (dpc), followed by the paired exoccipitals and basisphenoid. The presumptive sphenopalatine synchondrosis was found at 15 dpc; by 16 dpc, the perichondrium around the cranial base segments was interrupted at sites where synchondroses were forming. The presphenoid formed between 16 and 17 dpc. On 18 dpc, the entire presumptive cranial base was cartilaginous, and there was marked hypertrophy of chondrocytes and their lacunae in the areas of impeding basioccipital, basisphenoid and presphenoid bone formation. Cellular hypertrophy was the most advanced in the basisphenoid region, confirming the caudal-rostral sequence of cartilage maturation [52]. Throughout the first week of postnatal development, growth and ossification of the cranial base accelerated and the otic capsule ossified rapidly. By the fourth day after birth, the basioccipital-basisphenoid and basisphenoid-presphenoid synchondroses had reached full development, in analogy to mirror-image epiphyses [49].

DIFFERENTIAL GROWTH IN THE CRANIAL BASE SYNCHONDROSES

There is no reason why growth in the zones on either side of the growth center should be the same. In human fetuses, more bone deposition occurs on the ethmoid at the sphenoid synchondrosis and on the occipital bone at the sphenop-occipital junction. The sphenoid bone, therefore, shows less growth anteriorly and posteriorly. The increasing sella-foramen cecum distance after 2 years of age could mean, however, that the front part of the sphenoid grows faster postnatally than the ethmoid, and continues growing after the ethmoid has ceased [15] and the constancy of sella-basiphenoid synchondrosal distance could imply a continuing preponderance of growth on the sphenoid [53]. Enlow [54] suggested that growth may occur at different rates and to different extents on opposite sides of the plate, such that in the sphenop-occipital synchondrosis, the greater part of this lengthening occurs on the occipital side. It has been argued that these observations are better explained by posterior drift of the sphenoid rather than by inequalities of synchondrosal growth. Melsen [55] did not find any asymmetry in histological examinations of the human postmortem material. Hoyte [56] studied the basioccipital synchondroses in pigs and rabbits, by injecting live animals with alizarin red and sacrificing them at different intervals; the results demonstrated that growth in the basioccipital synchondroses is symmetric at each site but vary in patterned sequence between sites. Experiments using [3H]-thymidine radioisotope labeling in growing rats found a caudo-rostral gradient of growth rates in the cranial base cartilages, with the highest rates in the basioccipital and lowest in the rostral edge of the presphenoid. In other words, the posterior part of the sphenop-occipital synchondrosis has a greater amount of bone formation in its inferior part than its superior part, which leads to angular changes in the cranial base [45, 46, 57].

CRANIAL BASE SYNCHONDROSES AS GROWTH CENTERS

An area of hypothetical debate concerns synchondrosis growth capability, and whether it is independent from or dependent on other factors. As a remnant from the fetal chondrocranium, it should be an active and expansive growth center owing to its tissue separating power. Scott [58] hypothesized that cranial synchondroses by virtue of their interstitial growth, turbidity and resistance to deforming forces can separate growing bones at sutures, just as the brain or an eyeball can. On the other hand, others [27, 59] have suggested that cranial base synchondroses do not possess a growth potential of their own and are just show a mechanically obligatory response to the growth of other organs, such as the brain, and to the growth of organ spaces, such as the oral and nasopharyngeal spaces. According to this view, the growth of the craniofacial skeleton occurs as a secondary or adaptive response to its functional matrix. In the living body, load or muscular activity serves as an extracellular stimulus that is transmitted to cells and modulates their genetic growth and differentiation, leading to what is known as the functional matrix hypothesis [60].

In order to test these hypotheses, many transplant-implants studies have been performed. Petrovic and Charlier [61] showed that chondrogenesis continued in 8-day-old explants of the sphenop-occipital synchondrosis of the rat. Servoss [62] used [3H]-thymidine labeling in vivo and in vitro in mice, and found that in vitro sphenop-occipital synchondrosis not only continued to grow, but did so at a rate comparable to that of the in vivo counterparts. Kuroda et al. [63], repeating similar experiments in rabbits, noted that the sphenop-occipital synchondrosis maintained its structure and grew independently, but less than in vivo. These experiments indicate that cranial base synchondroses are endowed with an independent growth potential and are comparable to epiphyseal plates and the costochondral junction with regards their tissue-separating capacity [64, 65]. The independent growth potential of synchondroses [10] are also evident from the observation that their growth continues for a longer duration than that of the brain, which means the growth of the cranial base and the brain is not closely interdependent [19, 66]. The studies overall suggest that synchondroses are genetically coded in craniofacial growth and will develop even in the absence of functional activity. However, skeletal growth and the preservation of normal structure are modulated by environmental (epigenetic) factors, such as mechanical force, hormones and other growth factors.

TIME OF CLOSURE OF THE SPHENOP-OCIPITAL SYNCHONDROSIS

Controversies have arisen with regard to the closure time of the sphenop-occipital synchondrosis [34, 67, 68]. Ford [69] investigated human dry skulls and first reported that the closure time of the sphenop-occipital synchondrosis occurred...
late in the growth period between 20 and 25 years or between 17 and 20 years. A radiographic study by Irwin [70] and a tomographic study by Powell and Brodie [71] led to suggestions that incipient closure may already be discerned at the beginning of puberty. Melsen [4] reported that closure had taken place in the interval between eruption of the second and third molars. Computed tomography of the sphenoid-occipital synchondrosis showed signs of early fusion by age 8 years of age; the age of closure was 16 years in girls and 18 years in boys [72]. On the other hand, in postmortem histologic and micro-radiographic studies of the sphenoid-occipital synchondrosis, Ingerval and Thilander observed closure as the appearance of bony bridges and demonstrated that the average age of closure in females is around 14 years and that the sphenoid-occipital synchondrosis is never completely open in many girls older than 13 years and 9 months; the corresponding age for the boys is 16 years [3, 21].

The initial establishment of an osseous bridge between the adjoining bones in humans was investigated by Melsen [55] with human autopsy materials: the process occurs between the pubertal growth maximum and the cessation of sutural growth, which corresponds to the result from Thilander and Ingervall [21]. Latham [73] noted a rapid linear increase of the basioccipital length until the age of 9 years, the beginning of a plateau at about 10 years and then another, but very slow, increase until 18 years, indicating that there is a sequence of slower growth in the synchondrosis between 8 and 11 years. Bunny bridges first appear by about 12 years (females) and 13 years (males), and synostosis (all cartilage replaced by bone) takes place by the age of 16 to 17 years.

**ROLE OF SYNCHONDROSES IN GROWTH**

The cranial base forms the platform to which the rest of the skull grows and attaches. It provides and protects the crucial foramina through which the brain connects the face and the rest of the body [24]. As the basicranium grows, it elongates and flexes in the sphen-ethmoid, mid-sphenoid, and sphen-occipital synchondrosis [19]. Increases in basicranial breadth and length also occur in sutures (e.g., the occipito-mastoid), and the endocranial fossae of the basicranium deepens through drift, during which bone is resorbed and deposited along the superior and inferior surfaces, respectively [74]. This integrated growth occurs through many mechanisms, the most important of which are sutural expansion, synchondrosoal deposition, drift and flexion. As the brain expands, it generates tension along the endocranial surface of the neurocranial cavity, thereby activating osteoblast deposition within the intra-sutural periosteum throughout the upper portion of the vault, as well as drifting in the lower portions of the vault and cranial base and endochondral growth within certain synchondroses [23, 75].

In humans, facial growth is about 95% completed by 16 to 18 years, at least 10 years after the majority of the neuro-basicranial complex has reached adult size [76]. The basicranium and neurocranium, however, may have some influence on the growth of certain facial dimensions because the upper face articulates with the anterior cranial base and the anterior cranial fossa, while the mid-face articulates with the middle cranial fossa. In particular, the upper and middle portions of the face grow primarily by lateral drift and anterior displacement around the ethmoid and in front of the sphenoid [74, 77].

**SYNCHONDROSIS AND CRANIAL BASE ANGLE**

Cranial base flexion is a unique cranial feature of modern human beings and also a reflection of brain evolution. Compared with the cranial of quadrupeds, the cranial base angle in humans is relatively small. This difference is believed to be mainly due to the upright posture of humans and the increase in brain volume as well as frontal positioning of the eyes [29]. Fusion along the sphenoid-occipital synchondrosis is believed to be responsible for cranial base flexion, which develops in concert with the development of the upper airway and the ability to vocalize [78, 79]. During development, the anterior and posterior cranial base flexes at the sella turcica in the middle sagittal plane and thus constitutes an angle in the cranial base, termed the cranial base angle or saddle angle [10].

There are two well-known ontogenic models regarding cranial base flexion. Enlow [80] proposed that cranial base flexion during human prenatal and postnatal ontogeny is due to increased brain growth, relative to slower growth of the midline basicranium. This view is consistent with human fetal studies showing that growth along the cranial base is significantly slower than in other parts of the skull [81] and that fetal brain growth is markedly rapid at the same time [82]. Taken together, these findings imply that brain size increases relatively during human fetal life. The other ontogenic model, proposed by Ross and Ravosa and known as general spatial-packing, suggests that increases in brain size (volume) relative to length of the midline cranial base create a spatial packing problem that drives cranial base flexion and coronal reorientation of the petrous bones [83]. These spatial-packing hypotheses have been substantiated by numerous studies but not by others, owing to a basicranium that is less flexed than expected in the modern human brain [23, 84]. The alternative explanation to this problem is that the cranial base flexion is probably an intrinsic feature of human beings that is predominantly genetically determined. However, this hypothesis has not yet been supported by research [10].

A cranial base angle is already established in the chondrocranium between the prechordal and parachordal region of the cranial base in the human fetus. It has been generally presumed that the base flattens in the early stages of development by an increasing angle between the prechordal and parachordal portions and that this process later continues between the ossified part of the basicranium [85]. On the other hand, Diewert found little change during 7 weeks and 10 weeks in human cranial base angulations, while the head lifted up and back and the face developed. Thus, the cranial base angle is similar to that in later prenatal and postnatal stages, suggesting that distinctively human patterns develop early in the embryonic period [86].

Björk [1] reported that the cranial base angle gradually bends in the first years of childhood up to about 10 years, but others have shown that bending probably happens earlier in life, perhaps by 5 to 6 years [87] or even 2 to 3 years [88]. Differences in these measurements probably relate to
different baseline measurements and, perhaps much more importantly, also the internal aspect of the cranial base, which remains stable from early childhood onwards [29].

The cranial base angle (Fig. 2) is measured at three points antero-posteriorly and can be easily identified from cephalometric images; the most commonly used cephalometric landmarks are the nasion, sella turcica and basion. This measurement consists of two legs. The anterior leg, where the maxilla is attached, extends from the sella turcica (S) to the frontal-nasal suture (N). The posterior leg, where the mandible is attached, extends from the sella turcica (S) to the anterior border of the foramen magnum, defined as the basion (Ba). Therefore, any change in the cranial base angle can affect the relationships of the maxilla and mandible and influence the type of malocclusion. The posterior leg (BaS) of the cranial base angle (BaSN) may be tipped anteriorly or posteriorly, whereas the anterior leg (SN) may also be tipped up or down anteriorly by variation in either S or N vertically. Furthermore, variable lengths may compensate for any cranial deflection, such as an acute posterior leg that places the mandible forward, and this action can be negated by a long posterior leg that places both the basion and mandible posteriorly and vice versa [89].

![Diagram showing the cranial base angle (θ), the sphenoid (SOS) and the jaw relationship.](image)

The correlation between the cranial base angle and orthodontics was noticed and described in 1948 followed by Björk, Moss and Rickets in 1955 [91]. The assessment of orthodontic anteroposterior skeletal discrepancy of the jaws and arches should also account for the role of the cranial base [92]. The relationship of the cranial base with jaw position and classification of malocclusion has subsequently been examined extensively. Scott [93] further proposed one of the main factors that influence facial prognathism is the opening of the cranial base angle. This angle has a large individual variation, with a 5° standard deviation, and is markedly decreased in value from birth until the first year. It is also considered one of the craniofacial constants as it changes little during growth from 5 to 15 years [87, 94, 95]. Smaller linear and angular dimensions have been shown in class III patients, whereas class II subjects demonstrate an increased cranial base angle that leads to a more posterior position of the mandible [92, 96, 97]. The anterior cranial base and the middle cranial fossa have also been reported to be longer in individuals with class II malocclusion [98].

However, many studies have shown that the etiological relationship between the cranial base flexion and the type of malocclusion is not well supported [89, 99, 100]. It therefore appears that the cranial base angle is not a pivotal factor in determining malocclusion. Rather, the differential growth pattern and direction between the anterior cranial base and posterior cranial base imply that length and inclination of the cranial base are controlling factors of jaw position. Abnormalities of the posterior cranial base are related to mandibular prognathism, and those of the anterior cranial base to retrusive maxilla. These relationships are probably explained by the glenoid fossa’s location in the posterior cranial base; an elongated cranial base would bring the glenoid fossa back and the mandible with it [1, 93]. A Norwegian study has recently demonstrated differential growth between the anterior and posterior cranial base: in the male group, the growth curve of the anterior cranial base flattened out after 18 years but the posterior part increased until 21 years; no such difference was detected in the female group, where both growth curves flattened out after 15 years [101].

Other cephalometric characteristics resulting from skeletal type might influence the sagittal jaw position and relationship. For example, the flattening of the saddle angle is associated with a decrease of the posterior cranial base angle N-Ba-S, an elongation of the posterior part of the cranial base at Ba-point, and an enlarged distance N-Ar, all favoring an increased sagittal discrepancy of the jaws [91, 94]. A compensating mechanism associated with cranial base flexure was first described by Anderson and Popovich [102] in a study showing that the angle between the posterior cranial base and the ramus of the mandible closes in a highly correlated way to compensate for the opening of the cranial base flexure. This phenomenon tends to maintain the angle between the ramus and the anterior cranial base.

Therefore,
ABNORMALITIES IN GROWTH AND DEVELOPMENT

The sphenoid-occipital synchondrosis is an important link between the development of the cranial vault and that of the facial skeleton [1, 16, 26]. Premature growth arrest of the cranial base is widely regarded as the primary abnormality in syndromic forms of craniosynostosis [103]. In Klinefelter syndrome and Williams syndrome, the major craniofacial changes are located in the cranial base [104, 105]. In Cretinism syndrome, Turner syndrome [106] and Down’s syndrome [107], the craniofacial structure is characterized by a short retrognathic face owing to reduced cranial base length and increased angulations. In complete clefts of lip and palate, cranial base deviations in dimension and shape have been widely described. The data indicate that cleft lip and palate are not isolated malformations localized to the jaws, but malformations that also involve the cartilaginous cranial base [108, 109].

RECENT ADVANCEMENTS IN CRANIAL BASE RESEARCH

Significant advances in the study of craniofacial development have revealed the genetic and environmental basis of numerous craniofacial malformations, some of which do not even appear to be genetic in origin. This finding significantly deepens our understanding toward the fundamental principles of development, how genes regulate cell activity and thus mechanisms of determining the pattern and form of the craniofacial complex. Without the knowledge of gene activity and signaling transduction pathways, the elucidation of mechanisms that control development would be impossible.

Revelations of fibroblast growth factor receptor (FGFR), Sonic Hedgehog (Shh), SOX 9 and Cbfa1 genes are some examples of the genes that have been found to play critical roles in craniofacial development. SOX 9 has been found to be a master transcriptional factor that regulates early differentiation of mesenchymal cells into the chondrocyte lineage [110]. The absence of SOX 9 results in a severe skeletal dwarfism known as campomelic dysplasia [111]. Cbfa1/run2/Osf2 serves as the earliest transcriptional regulator of osteoblast differentiation and can be considered to be the central regulator of intramembranous ossification [112]. Mutations involving Cbfa1 result in cleidocranial dysplasia in humans [5]. Shh, a member of the vertebrate Hedgehog gene family, plays an important role in establishing craniofacial midline structures and in regulating cranial suture development [113]. Shh is the most commonly identified single-gene defect causing human holoprosencephaly [114]. Mutations in genes encoding FGFRs have been identified as causes of Apert syndrome, Crouzon syndromes and Pfeiffer syndromes [5]. As for environmental effects, the use of an organ culture technique has revealed that synchondrosis is responsive to tensile mechanical stress and increased expression of various factors related to growth stimulation [115-117]. This series of studies was the first to show that growth across the synchondrosis can be modified mechanically.

CONCLUSION

Cranial base synchondroses, with their unique bipolar growth, are a fundamental part of craniofacial development. Abnormalities in their development have serious consequences because of their position and growth in the craniofacial complex. Molecular biology and genetics provide tools to further investigate the molecular mechanisms, genes and transcription factors responsible for synchondrosis and craniofacial development.

REFERENCES


Relationships Between Cranial Base Synchondroses and Craniofacial Development

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