

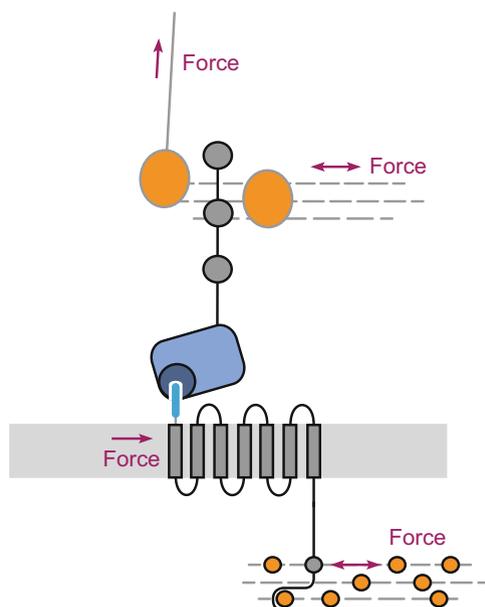
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# Adhesion GPCRs as a Putative Class of Metabotropic Mechanosensors

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## Graphical Abstract



Adhesion GPCRs as mechanosensors. Different aGPCR homologs and their cognate ligands have been described in settings, which suggest that they function in a mechanosensory capacity. For details, see text

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### Abstract

G protein-coupled receptors (GPCRs) constitute the most versatile superfamily of biosensors. This group of receptors is formed by hundreds of GPCRs, each of which is tuned to the perception of a specific set of stimuli a cell may encounter emanating from the outside world or from internal sources. Most GPCRs are receptive for chemical compounds such as peptides, proteins, lipids, nucleotides, sugars, and other organic compounds, and this capacity is utilized in several sensory organs to initiate visual, olfactory, gustatory, or endocrine signals. In contrast, GPCRs have only anecdotally been implicated in the perception of mechanical stimuli. Recent studies, however, show that the family of adhesion GPCRs (aGPCRs), which represents a large panel of over 30 homologs within the GPCR superfamily, displays molecular design and expression patterns that are compatible with receptivity toward mechanical cues (Fig. 1). Here, we review physiological and molecular principles of established mechanosensors, discuss their relevance for current research of the mechanosensory function of aGPCRs, and survey the current state of knowledge on aGPCRs as mechanosensing molecules.

### Keywords

Adhesion GPCR • Mechanosensor • Mechanosensation

## 1 General Overview of Mechanosensation

Each cell in our body is constantly exposed to a multitude of mechanical cues emanating from cell movements, hydrostatic pressure, shear stress exerted by fluids, or compressive and tensile forces of various origins. In order to react to mechanical conditions, these need to be (1) conveyed to mechanosensitive

molecules (a process termed *mechanotransmission*; Fig. 2). (2) These molecular mechanosensors perceive force application through conformational changes (*mechanoreception* or *mechanosensing*). (3) Force-dependent changes in mechanosensor structure are finally handed over to intracellular signaling cascades that are not necessarily force dependent per se, including metabotropic effectors and transcriptional pathways, which finally shape the cellular response to the mechanical change (*mechanoreponse*) [1]. Here, we will concentrate on the role of aGPCRs in the process of mechanoreception.

Mechanosensors frequently constitute integral membrane proteins, and it is commonly accepted that mechanical forces that impinge onto the cell surface affect local and global membrane tension. Thus far, a number of molecules have been attributed with a function in mechanoreception, among them ion channels, cell surface receptors (e.g., integrins, cadherins/selectins), and, more recently, G protein-coupled receptors (GPCRs).

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## 2 Established Mechanosensors

Mechanosensors evolved different receptive strategies to reliably perceive, integrate, and convey mechanical information into the cell. Even though detailed molecular knowledge of the mechanotransduction cascade is, for many mechanosensors, still unknown, an understanding of the receptive strategies of these molecules will help define to what extent aGPCRs serve as metabotropic mechanosensitive devices. Thus, the following section delineates potential mechanisms that underlie mechanotransduction and provides an overview of established mechanosensors.

### 2.1 Molecular Models of Mechanosensation

Two models currently encompass how mechanical strain is transmitted onto mechanosensors. The “tethered” model is based on the concept that surface molecules are anchored to intra- and/or extracellular components via molecular springs, which convey mechanical forces to membrane integral mechanosensors (e.g., TMC, extracellular; see Sect. 2.2.5 NOMPC, intracellular; see Sect. 2.2.3). Alternatively, in the “membrane” model, mechanosensitivity is shaped by the lateral pressure profile [2] at the receptor/bilayer interface promoting conformational changes of mechanoreceptive proteins. Given that a cell’s response to force is signified by its mechanical context by means of the surrounding extracellular matrix (ECM), neighboring cells, and the mechanical properties of the cell itself, it is reasonable to assume that both mechanisms are employed in parallel to ensure reliable integration of physical forces.

## 2.2 Ionotropic Mechanosensors

Guharay and Sachs were the first to discover that skeletal muscles in chick express an ion channel that can be activated by membrane stretch [3, 4]. After that, several years passed until the first mechanosensitive channel genes, *mscL* and *mscS*, were identified and cloned from prokaryotes [5–7]. In *Bacteria* and *Archaea*, mechanosensitive channels are primarily involved in detecting osmotically induced cell swelling to regulate turgor [6, 8].

A large number of mechanosensitive channels were identified in eukaryotic cells [9] and, based on their ion permeability, can be divided into two classes: excitatory cation-selective ion channels and inhibitory depolarization-gated ion channels. We will briefly discuss examples for both groups below. Furthermore, a novel ion channel family—transmembrane channel-like 1 and 2 (TMC1/TMC2)—was recently identified.

### 2.2.1 Piezo

Vertebrate Piezo1 and Piezo2 (or Fam38A and Fam38B) channel subunits comprise large membrane proteins with 14 predicted membrane-spanning segments [10]. Heterologous expression of Piezo conveys mechanosensitivity to a mechanically insensitive cell [11]. Furthermore, functional ion channels could be reconstituted from Piezo proteins in liposomes [12]. Thus, ectopically expressed Piezo appears to form mechanosensitive ion channels independent from cytoskeletal elements implying that the channel activating tension originates from the membrane [11–13].

Mammalian *Piezo 1/Piezo 2* was detected in the lung, bladder, colon, and skin. *Piezo 2* was additionally enriched in dorsal root ganglia (DRGs) [11], where it acts as a mechanotransducer to regulate the perception of tactile stimuli [14, 15]. Similarly, the *piezo2b* homolog plays a role in zebrafish touch sensation [16]. Moreover, Piezo1 was assigned roles in the regulation of erythrocyte cell volume and vascular development in zebrafish and mouse, respectively [17–19]. The sole Piezo homolog in *Drosophila* (*dmpiezo*) produces, similar to its vertebrate counterpart, mechano-dependent ion currents and is required for the perception of noxious mechanical stimuli *in vivo*. Interestingly, Piezo appears to act in parallel to Pickpocket (PPK), which belongs to the (DEG)/ENaC family, to regulate mechanical nociception in *Drosophila* larvae [20].

### 2.2.2 DEG/ENaC Channel

The DEG/ENaC channel family was named after their first members, degenerins (*mec-4* and *deg-1*) and epithelial sodium channel (ENaC) from *C. elegans* and mammals, respectively [21–24]. DEG/ENaC channels are expressed in different tissues across a large range of phyla, are usually Na<sup>+</sup>-selective, and appear to be activated by a wide spectrum of stimuli including mechanical force [25]. Two transmembrane domains (N- and C-termini intracellular) connected by a large extracellular loop define the protomer architecture of DEG/ENaC proteins, which can form both homo- and heteromeric ion channels [26–28].

The first identified eukaryotic genes involved in touch sensation, *mec-4* and *mec-10* (mechanosensory abnormal) [29], were uncovered through forward genetic screens in *C. elegans*, in which mutants were probed for their ability to respond to gentle body touch conveyed through a defined set of touch receptor neurons [29–31]. Genetic, biochemical, functional, and morphological analyses implicate four MEC proteins (MEC-4, MEC-10, MEC-2, MEC-6) to compose mechanosensory channel complexes [32–34]. The pore-forming subunits are presumably contributed by MEC-4 and MEC-10 [23, 32, 35, 36]. Interestingly, the local lipid milieu appears to influence the function of this mechanosensory complex [33, 37–39], suggestive of a membrane stretch-mediated gating mechanism.

### 2.2.3 TRP Channels

Transient receptor potential (TRP) channels are involved in recognizing a broad range of mechanical and chemical stimuli [40]. Each TRP channel gene belongs to one of seven subfamilies [TRPC (classical), TRPV (vanilloid), TRPM (melastatin), TRPN (NOMPC), TRPA (ANKTM1), TRPML (mucoipin), and TRPP (polycystin)] [41] and encodes a channel subunit with at least six transmembrane-spanning segments and intracellular N- and C-terminal regions with subfamily-specific domain layout [42]. Four subunits assemble into functional homo- or heteromeric ion channels, commonly nonselective for cations and permeable to  $\text{Ca}^{2+}$  [41, 43].

Mechanosensitivity has been reported for a number of TRP channel subunits; however, to what extent these proteins are directly involved in mechanotransduction remains to be resolved for many family members. Interestingly, TRP channels were also identified as partners of GPCRs and aGPCRs in this context.

For example, TRPC6 is a downstream target of a mechanosensitive GPCR responsible for mediating myogenic vasoconstriction in response to elevated intraluminal pressure [44]. Further, mouse sensory neurons express TRPC1, TRPC3, and TRPC6 [45]. *Trpc3/Trpc6* double knockout mice display altered responses to tactile stimuli. These mutants also exhibit pronounced hearing impairments and vestibular defects consistent with TRPC3 and TRPC6 expression in cochlear and vestibular hair cells [45]. Recent work indicated that TRPC1, TRPC3, TRPC6, and TRPC5 heteromultimerize to contribute to cutaneous and auditory mechanosensation. Moreover, these TRP channel subunits were proposed to contribute indirectly to cochlear mechanotransduction [46].

Further, members of the TRPN family have been associated with the perception of mechanical stimuli in *Drosophila* (no receptor potential; NOMPC) [47, 48], hearing in zebrafish (TRPN1) [49], and proprioception in *C. elegans* (TRP-4) [50]. NOMPC localizes to the ciliary tips of mechanosensory neurons of the Johnston's organ and chordotonal organs in *Drosophila* [51, 52]. NOMPC can be directly mechanically activated and confers mechanosensitivity to otherwise insensitive cells, rendering NOMPC a bona fide mechanotransduction channel [53, 54]. Recent data indicate that the 29 ankyrin repeats of NOMPC's N-terminus constitute a molecular spring that conveys forces generated by the

cytoskeleton to modulate ion channel gating [55, 56]. However, loss of *nompC* does not completely abolish fly hearing [47], suggesting the existence of additional transducer molecules [57]. Alternatively, NOMPC may sensitize or adjust the mechanosensory complex for mechanical input [58]. Interestingly, genetic analysis suggests that the aGPCR dCIRL/latrophilin modulates NOMPC activity in sensory chordotonal neurons [59] (see Sect. 3.2).

The *C. elegans* TRPV proteins OSM-9 and OCR-2 form heteromeric channel complexes in sensory neurons that respond to touch and hyperosmolarity [60–62]. Consequently, loss of function of *osm-9* results in worms that are less sensitive to osmotic and mechanical challenges [60, 62]. *Drosophila* encodes TRPV proteins, INACTIVE and NANCHUNG, that are mutually required for the assembly of heteromultimeric channels involved in fly hearing and possibly auditory mechanotransduction [57, 58, 63, 64]. Epistatic analysis implicated NANCHUNG function in a mechanosensory signaling pathway together with the aGPCR latrophilin/CIRL [59].

Finally, TRPP homologs TRPP2 (polycystin 2 or PC2) and TRPP3 (polycystic kidney disease 2-like 1, PKD2L2) have been associated with mechanosensation. TRPP2 forms a receptor-ion channel complex with polycystin 1 (PC1, TRPP1) [65, 66], which localizes to primary cilia of renal epithelial cells and endothelial cells [67–70]. PC1 does not belong to the TRP channel family; it contains 11 - transmembrane-spanning regions and a large extracellular N-terminal domain that promotes cell-cell and cell-matrix interactions [71, 72] (see also [73]). The TRPP2/PC1 complex mediates  $\text{Ca}^{2+}$  transients in response to ciliary deflections induced by luminal shear stress [67, 68, 74]. Intriguingly, filamin A cross-links TRPP2 and the actin cytoskeleton to regulate stretch-activated cation channels (SACs) involved in cardiovascular pressure sensing [70]. Furthermore, TRPP2 forms ciliary mechanosensitive sensors with TRPV4 to transduce mechanical stress [75].

#### 2.2.4 $\text{K}_{2\text{P}}$ Channels

$\text{K}_{2\text{P}}$  channels are comprised of four transmembrane-passing segments, two pore-forming regions, and intracellular N- and C-terminal portions. They assemble into hetero- and homodimeric ion channels. TWIK-related  $\text{K}^+$  channels (TREK) and TWIK-related arachidonic acid-stimulated  $\text{K}^+$  channel (TRAAK) are membrane tension-gated channels characterized by low mechanical threshold and a broad range of tension activation [76].

TREK-1 and TREK-2 are widely expressed within the central and peripheral nervous systems as well as nonneuronal tissues such as the cardiovascular system, lung, colon, and kidney. By contrast, TRAAK expression appears to be confined to neuronal tissues [77]. In the nervous system,  $\text{K}_{2\text{P}}$  channels are physiologically relevant as they contribute considerably to cell hyperpolarization, which balances/counteracts depolarization-induced action potentials, thereby shaping the electrical response to mechanical force. For example, TRAAK activity counteracts activation of Piezo to curtail action current generation in cultured neuroblastoma cells [78]. Further, TRAAK/TREK and mechanosensitive cation channels are coexpressed in sensory neurons of DRGs [79]. Deletion of *Traak*, *Trek-1*, or

*Trek-2* genes in mice renders them hypersensitive to mechanical challenge [80, 81], which is in line with the notion that  $K_{2P}$  channels set the mechanical threshold for action potential generation [76]. Evidence for the physiological relevance of TREK-1 in mechanosensation outside the nervous system has emerged as well. Polycystins play a role in force-dependent apoptosis of renal epithelial cells, a function that was shown to rely on the opening of TREK-1 [82]. TREK-1 is enriched in the bladder, uterus, and colon where it regulates stretch-induced contraction of smooth muscle cells [83].

### 2.2.5 Transmembrane Channel-Like 1 and 2 Proteins

Recently, the transmembrane channel-like 1 and 2 (TMC1 and TMC2) proteins emerged as novel components required for auditory and vestibular mechanosensation in mammals [84–86]. The *Tmc* gene family encodes proteins with at least six transmembrane regions flanked by intracellular N- and C-terminal portions [87] reminiscent of TRP channel proteins. However, TMC proteins do not share sequence homologies with known ion channels [85, 88], even though recent work indicated that *C. elegans tmc-1* assembles into nonselective cation channels when heterologously expressed [89]. Several lines of evidence implicate TMC proteins as key components of the hair cell transduction complex. First, consistent with the mRNA expression pattern [90], TMC1 and TMC2 fusion proteins localize to stereocilia of the inner ear hair cells [86]. Second, mutations of the human *Tmc1* gene have been associated with dominant and recessive nonsyndromic sensorineural hearing deficits [84]. Similarly, dominant and recessive mutations of mouse *Tmc1* result in animals that exhibit defective hearing (*Beethoven*) and deafness (*dn*), respectively [84, 91]. Third, in mice, loss of function of *Tmc1* or *Tmc2* deprives hair cells of mechanosensory responses [90, 92]. Fourth, robust apical FM1-43 dye or gentamicin uptake reported for wild-type hair cells was abolished in the absence of TMC1 and TMC2 [90, 93, 94]. Fifth, expression of either TMC1 or TMC2 rescues loss of hair cell mechanosensitivity displayed by homozygous null mutants [90]. Sixth, protocadherin-15 (PCDH15) together with cadherin-23 forms extracellular tethers (called tip links), which connect adjacent pairs of stereocilia and transmit hair cell bundle deflection-based mechanical forces to mechanosensors [95, 96]. Strikingly, recent work uncovered a direct interaction between PCDH15 and TMC1 as well as TMC2 in fish and mouse [97, 98].

Taken together, these results suggest that TMC proteins are strong candidates for the long sought-after mechanosensitive channel required for mechanotransduction in mammalian hair cells [85].

## 2.3 Integrins and Cadherins

Cell adhesion enables the generation of mechanical tissue cohesion as well as cell-cell and cell-matrix communication vital for a myriad of physiological processes. But how do cells sense the mechanical context of their microenvironment (e.g., extracellular matrix or adjacent cells), and how are these signals translated into

cellular responses? Based on structural and functional properties, integrins and cadherins have been associated with mechanosensation in this vein.

### 2.3.1 Integrins

Integrins constitute a large family of glycoprotein receptors that bridge cell-cell contacts (cell adhesion) and interconnect intra- and extracellular matrices (ECMs) in metazoans. Integrins exist as heterodimers comprised of non-covalently bound  $\alpha$ - and  $\beta$ -subunits, which are combined to form more than 20 distinct, functionally nonredundant receptor variants [99, 100].

Integrin heterodimers are inserted into the plasma membrane through a single transmembrane domain per protomer. The large extracellular domain (ECD) mediates interactions with ECM components (e.g., collagens, fibronectins, or laminins) and adjacent receptors [e.g., vascular cell adhesion molecule-1 (VCAM-1) and intercellular cell adhesion molecule (ICAM)] [100, 101]. The short intracellular domain (ICD) interacts, via cytoplasmic adaptors, with cytoskeletal elements to modulate local actin polymerization and global cytoskeletal dynamics [102, 103]. This molecular layout allows integrins to mechanically interlink the cell's exterior and interior providing the backbone for mechanical stress-induced assembly of focal adhesions (FA) [104–107], stimulation of signaling pathways, and gene transcription [108–110].

Importantly, integrins possess the capacity to confer signals in an “outside-in” or “inside-out” fashion associated with different biological ramifications. “Outside-in” signaling is based on the interaction of integrin ECDs with ECM molecules or opposing cellular receptors. These engagements provide information about ECM rigidity as well as adjacent cell/tissue geometry, which in turn regulate cellular growth, differentiation and apoptosis, cell polarity, and formation of focal adhesion complexes [105, 111–114]. During “inside-out” signaling, intracellular activators such as kindlins or talins associate with integrin ICDs to induce conformational changes that alter the affinity to extracellular ligands (“integrin activation”). Hence, this signaling mode affects cell adhesion and migration as well as ECM assembly [102]. Interestingly, the ionotropic Piezo1/Fam38A channel subunit was shown to induce integrin activation through an R-Ras-dependent mechanism [115].

In sum, integrin-based adhesion complexes are fit to perceive and integrate external forces into the preexisting cellular mechanical context to balance the ECM resistance according to tension forces generated by the cytoskeleton.

### 2.3.2 Cadherins

Cadherins are a large group of cell surface receptors that mediate  $\text{Ca}^{2+}$ -dependent cell adhesion [116]. The cadherin family includes a vast number of cell surface molecules; however, classical cadherins (E-, N-, P-cadherins) and the related desmosomal cadherins are, thus far, the best characterized. They feature cadherin repeat-containing ECDs, a single membrane-spanning region, and ICDs which interact with  $\beta$ -catenin (classical cadherins),  $\gamma$ -catenin (desmosomal cadherins), and components of the cytoskeleton [117].

Cadherins engage in ECD-dependent homo- and heterophilic interactions that may mediate adhesive and selectivity functions, respectively [118]. For example, force measurements uncovered that cadherins establish three spatially distinct bonds at different ECD regions that possess varying kinetic and mechanical properties [119, 120]. In contrast, ICDs associate with F-actin via catenin/vinculin interactions [121–124]. This structural setup allows cadherin complexes to act as mechanosensors that actively sense and transduce exogenous and endogenous mechanical fluctuations to trigger biochemical changes that direct cell responses [125]. Strikingly, the tensile force administered to an adhesion site is directly proportional to ECM stiffness [126].

Several lines of evidence support the role of cadherin-based adhesive junctions in mechanotransduction. First, cell-cell junctions undergo force-dependent remodeling not only *in vitro* but also *in vivo* [127, 128]. In *Drosophila* and *C. elegans*, cytoskeletal-provoked tension at cadherin junction coincides with a decline in junction expansion in various developmental processes. For example, the association of the actomyosin apparatus with cadherin junctions is a prerequisite for contraction-driven apical constriction during *C. elegans* gastrulation. A similar mechanism has been reported for ventral furrow formation in *Drosophila* [129]. Intriguingly, recent work demonstrated that during germ band extension in *Drosophila*, junction shrinkage appeared to result from tension-generated cadherin accumulation, which was counteracted by cadherin endocytosis [130]. Second, the mechanics of intercellular junctions (e.g., cell traction, adhesion strength, and junction rigidity) change according to the mechanical environment [122, 131–134].

In *Drosophila*, junctional mechanics are regulated in a forward loop that controls cadherin levels with respect to actomyosin density to enable adequate tissue morphogenesis [130, 135].

Taken together, cadherin complexes probe their environment for tensional changes to preserve the mechanical integrity of cellular junctions and to regulate morphogenesis and homeostasis [136–139].

## 2.4 Metabotropic Mechanosensors

GPCRs are biosensors vital for the transduction of light, olfactory and gustatory stimuli, as well as neurotransmitters and hormones into biochemical signals and thus shape a multitude of physiological processes [140]. Based on this perception profile, GPCRs were classically considered chemoreceptors. However, in recent years, mounting evidence suggests that mechanical forces can also be perceived through GPCRs. Therefore, GPCRs most likely constitute polymodal sensors with both chemoreceptive and mechanoreceptive properties.

The first mechanosensitive GPCR identified was the angiotensin II type 1 receptor (AT<sub>1</sub>R), which mediates functions in preload-induced cardiac hypertrophy *in vitro* and *in vivo*. In contrast to autocrine-mediated vasoconstriction, activity of AT<sub>1</sub>R is brought about independently of angiotensin II and facilitates intracellular signaling through G<sub>q/11</sub> proteins [141, 142]. In addition to this

pathophysiological function, intraluminal pressure stimulates AT<sub>1</sub>R<sub>s</sub> located in the arterial blood vessels causing myogenic vasoconstriction, a mechanism referred to as the Bayliss effect [44]. Mounting evidence implicates PLC kinases in mechano-dependent signaling pathways, which may subsequently trigger activation of ionotropic TRPC channels (see Sect. 2.2.3) to set arterial myogenic tone [44, 141, 143, 144].

Strikingly, AT<sub>1</sub>R undergoes mechanically induced conformational changes that stabilize the seventh transmembrane domain near the agonist-binding pocket, a molecular organization that is different from that induced by agonist and inverse agonist signaling [145–147], suggesting stimulus modality-dependent signaling modes of this GPCR.

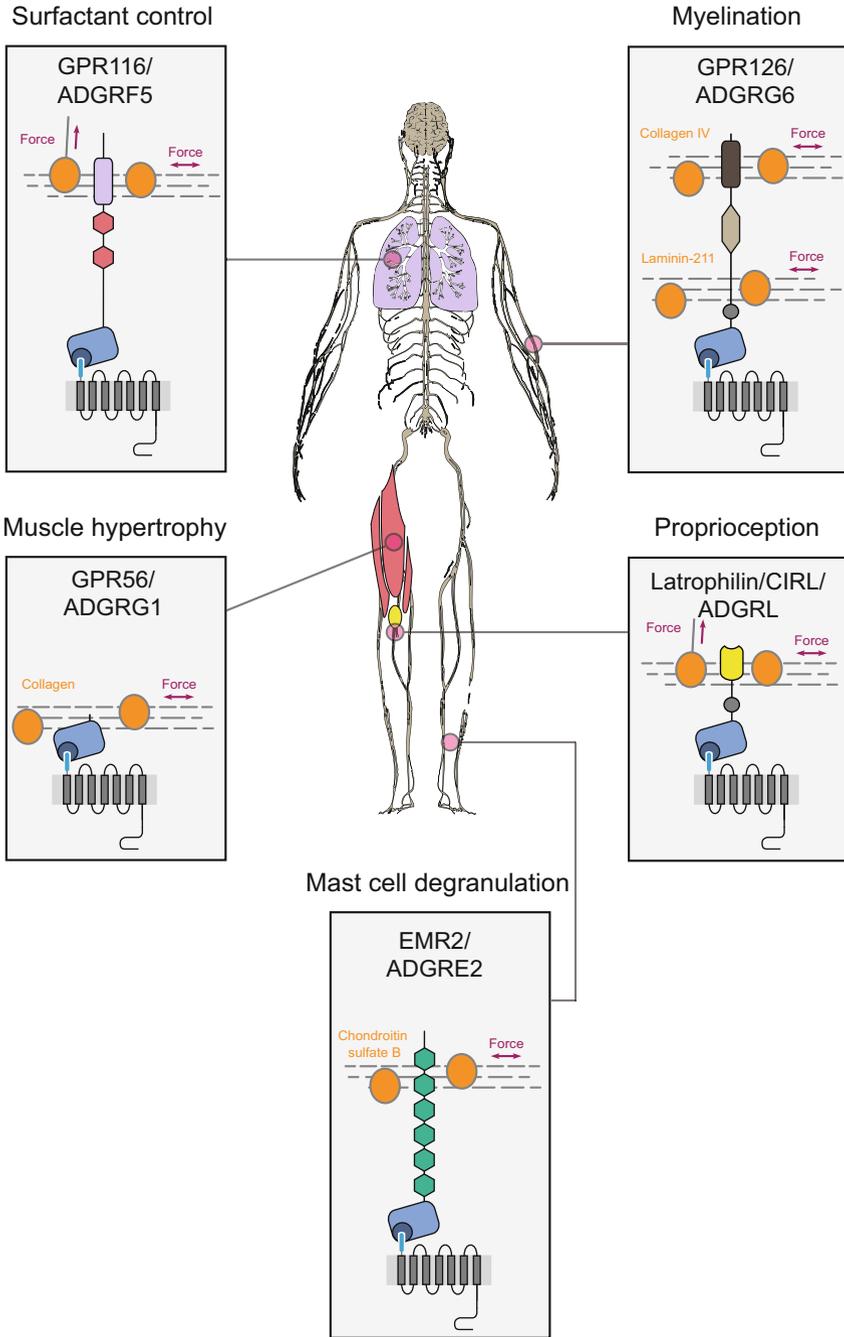
Thus far, eight additional putative mechanosensitive G<sub>q/11</sub> protein-coupled receptors have been described [148]. For example, time-resolved fluorescence microscopy of endothelial cells unveiled that mechanical strain causes a ligand-independent rise in bradykinin 2 receptor (B<sub>2</sub>R) activity [149]. The authors suggest that, similar to AT<sub>1</sub>R, conformational alterations of B<sub>2</sub>R are due to tension-based properties of the lipid bilayer. However, it remains unclear if mechanical stimulation of B<sub>2</sub>R is causally linked to effective G protein coupling. Another interesting example of a potential mechanosensitive GPCR derives from type 1 parathyroid hormone receptor (PTH<sub>1</sub>R) essential for Ca<sup>2+</sup> homeostasis in osteoblasts [150, 151]. Bone formation and bone mass regulation rely on the availability of parathyroid hormone (PTH) and mechanical loading, signal modalities proposed to converge at PTH<sub>1</sub>R [151]. Evidence for a role in mechanoreception was also reported for dopamine receptor type 5 (D<sub>5</sub>R) and formyl peptide receptor (FPR<sub>1</sub>), which are both, similar to PTH<sub>1</sub>R, excitable through fluid flow-generated shear stress in endothelial cells and neutrophils, respectively [152, 153]. Interestingly, D<sub>5</sub>R localizes to primary cilia *in vitro* and *in vivo* and regulates their length through cofilin and actin polymerization. In addition, the authors report that ciliary sensitivity to fluid shear stress can be modified through chemosensory properties of the receptor [152]. Furthermore, electrophysiological recordings unveiled hypotonicity-dependent responses of heterologously expressed vasopressor receptors, ET<sub>A</sub> endothelin receptor (ET<sub>A</sub>R) and V<sub>1A</sub> receptor (V<sub>1A</sub>R), as well as H<sub>1</sub> histamine receptor (H<sub>1</sub>R) and M<sub>5</sub> muscarinic acetylcholine receptor (M<sub>5</sub>R) [44].

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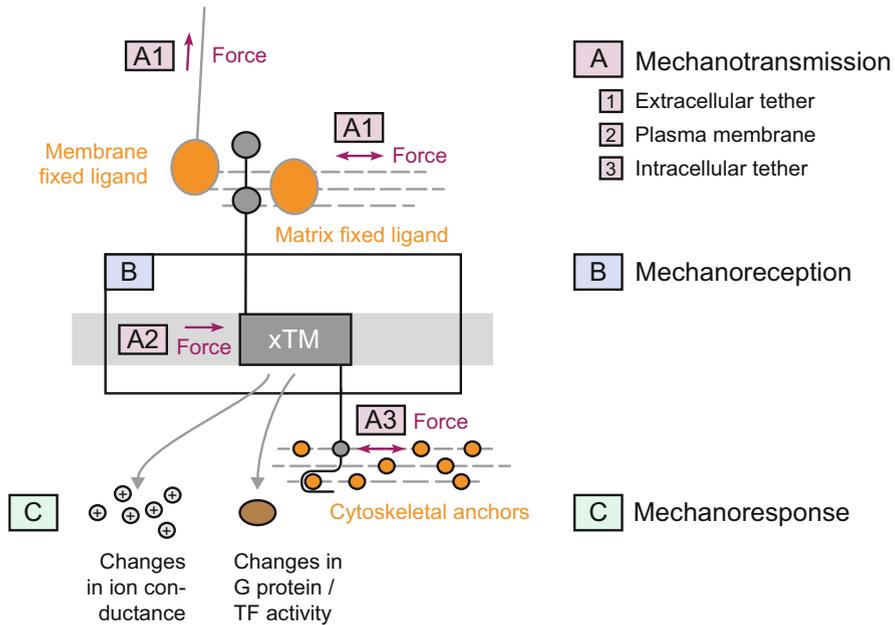
### 3 Adhesion GPCRs in Mechanosensation

A novel intriguing line of evidence for the mechanoreceptive role of GPCRs derives from the large family of aGPCRs, which stand out from canonical GPCRs because of their structural and functional profile as well as the conspicuous lack of soluble ligands [154]. Thus, are aGPCRs particularly prone to sense mechanical cues?

Three members of the aGPCR class, GPR56/ADGRG1, GPR126/ADGRG6, and latrophilin/CIRL/ADGRL, have been recently suggested to be involved in mechanosensation (Fig. 1) [59, 155, 156]. Like canonical GPCRs, aGPCRs possess a seven-transmembrane helix region (7TM) that can intracellularly couple to



**Fig. 1** Adhesion GPCRs as mechanosensors. Different aGPCR homologs and their cognate ligands have been described in settings, which suggest that they function in a mechanosensory capacity. For details, see text

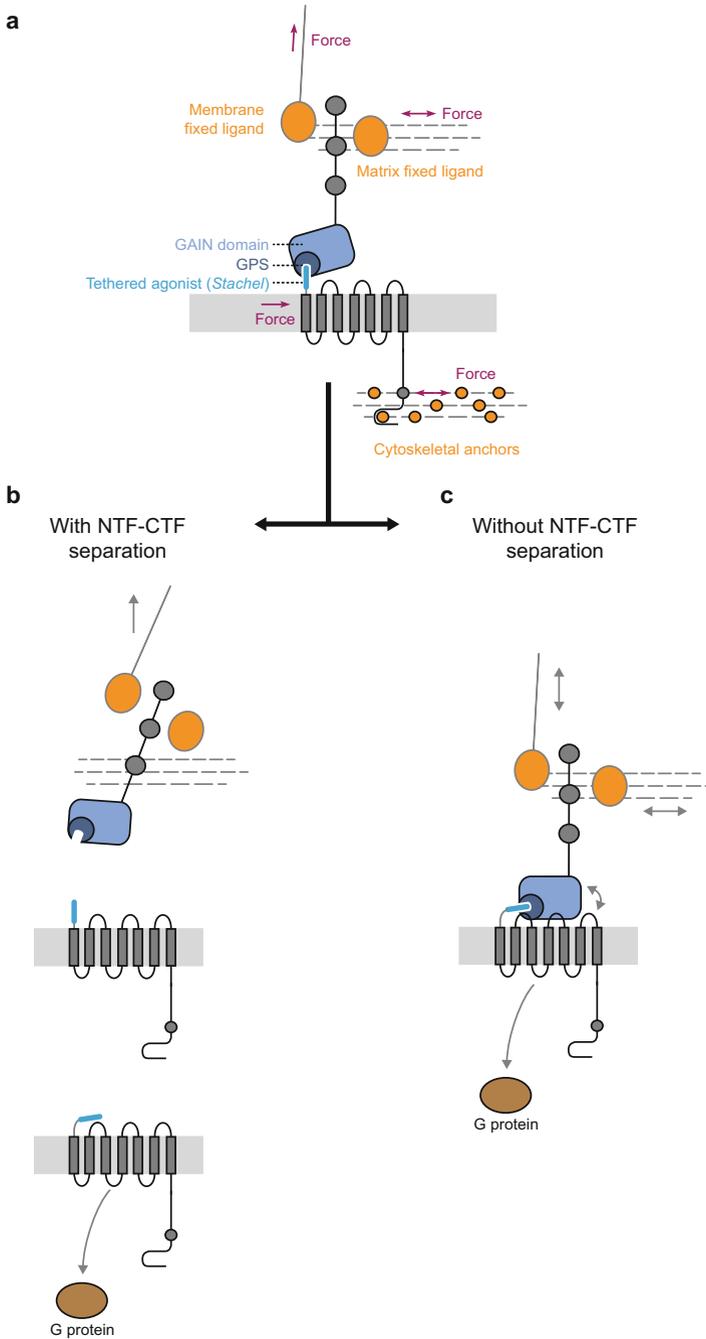


**Fig. 2** Principal elements of mechanotransduction. The process of mechanotransduction can be grossly divided into three steps. (a) Mechanotransmission (indicated in *light pink*) that, according to current models, can be conveyed through extracellular or intracellular tethers and/or the plasma membrane. (b) Mechanosensation (*blue*), perceived through force-dependent conformational changes of membrane integral sensor proteins (*gray*). (c) Mechanoresponse (*green*), which induces changes in cell physiology (e.g., ion channel conductances, metabolic states, or modulation of transcriptional activity). xTM = variable number of transmembrane helices depending on the mechanosensor protein architecture (TRP channels, 6TM; aGPCR, 7TM; polycystin 1, 11TM; Piezo, 14TM)

heterotrimeric G proteins capable of triggering a multitude of signaling cascades that determine cellular responses (Fig. 2; see [157, 158]).

Further, aGPCRs possess large extracellular N-termini, which for many aGPCRs contain domains that are involved in cell-cell or cell-matrix adhesion [154] (see also [159]). Uniquely, the N-termini of many aGPCRs can be separated from the 7TM through an autoproteolytic event catalyzed by the GPCR proteolytic site (GPS) motif [160, 161], which is part of a much larger protein fold (GPCR autoproteolysis-inducing (GAIN) domain) [162]. Receptor cleavage is thought to occur early in the secretory pathway during protein maturation and generates an N-terminal fragment (NTF) and C-terminal fragment (CTF), which are predicted to traffic together to the cell surface where they are located as non-covalently bound heterodimers (Fig. 3; see also [73, 159]).

For several aGPCRs, a cryptic tethered ligand, termed the *Stachel* sequence, has been described, which acts as a potent agonist for downstream signaling [163–166] (Fig. 3; see also [157, 158]). Yet, crystal structures of aGPCR GAIN domains



**Fig. 3** Sequence of putative events during aGPCR-mediated mechanotransduction. **(a)** aGPCRs are subjected to mechanical force through their adhesive ECDs, at the level of the membrane, or through intracellular anchors. **(b)** Depending on whether the receptor is autoproteolyzed and

demonstrate that the *Stachel* sequence is deeply buried between two  $\beta$ -sheets [162], suggesting that the tethered agonist, at least for some aGPCRs, may not be readily exposed. How, then, might the *Stachel* sequence “escape” the ensconcement of the GAIN domain  $\beta$ -sheets in order to confer a signal to the receptor’s 7TM stretch? Recent studies suggest that mechanical activation may be at play, which could induce conformational changes within the GAIN domain or complementary portions of the receptor to liberate the *Stachel* sequence and to allow receptor activation (Fig. 3).

### 3.1 GPR126/ADGRG6

As described in [167], myelin is the multilamellar sheath generated by specialized glial cells called Schwann cells in the peripheral nervous system and oligodendrocytes in the central nervous system. In the peripheral nervous system, GPR126 couples to  $G\alpha_s$  and elevates cAMP in Schwann cells to initiate myelination (Fig. 1) [155, 163, 168–171]. The *Stachel* sequence of GPR126 is a potent agonist to trigger cAMP elevation, and analysis of zebrafish mutants has recently provided evidence that *Stachel*-mediated signaling is critically important in vivo. Liebscher and colleagues generated a new mutant allele of *gpr126*, *gpr126<sup>stl215</sup>*, in which two amino acids essential for signaling were precisely deleted from the *Stachel* sequence. In vitro analysis demonstrated that this mutant receptor could be cleaved and that it trafficked appropriately to the cell surface, although it was incapable of signaling via cAMP. Accordingly, in vivo, *gpr126<sup>stl215</sup>* mutant Schwann cells could not initiate myelination [163]. This work suggested a model in which *Stachel*-mediated activation of Gpr126 is required for cAMP elevation and subsequent myelin initiation.

To determine how the *Stachel* sequence of GPR126 might be exposed during Schwann cell myelination, Petersen and colleagues investigated the relationship between one GPR126 binding partner, laminin-211, and GPR126 signaling [155] (discussed further in [167]). Laminin-211 is a heterotrimeric protein encoded by *Lama2*, *Lamb1*, and *Lamc1* genes [172], and like GPR126, laminin-211 is required for Schwann cell development and myelination [173]. Overexpression of *lama2* in zebrafish rescues myelination in *gpr126* hypomorphic mutants in a cAMP-dependent manner, suggesting that laminin-211 activates  $G\alpha_s$  signaling.

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**Fig. 3** (continued) present as an NTF/CTF heterodimer at the cell surface, force transmission may separate the NTF from the CTF, thereby exposing the cryptic tethered agonist that interacts with the 7TM domain and triggers intracellular responses. (c) Several aGPCRs cannot be autoproteolytically cleaved, yet they appear to harbor a *Stachel* sequence that can stimulate receptor activity. This indicates that the tethered agonist can interact with the 7TM domain even if it remains associated with GAIN domain, possibly through conformational changes of the GAIN domain that render the agonist accessible. Note that this scheme illustrates putative intramolecular interactions between GAIN/*Stachel* and the 7TM region of aGPCRs; however, intermolecular interactions are conceivable as well

Intriguingly, addition of laminin-211 to GPR126 in heterologous cell systems suppressed cAMP elevation under standard culture conditions and elevated cAMP levels under dynamic conditions (vibration or shaking) [155]. These data can be reconciled by observations from related aGPCRs, noted above, that the *Stachel* sequence is buried, not exposed, between two  $\beta$ -sheets of the GAIN domain [162]. The *Stachel* sequence is so deeply buried that significant structural changes might be required for *Stachel*-mediated receptor activation to occur (Fig. 3). In vitro analyses of mutant receptors demonstrated that laminin-211 addition facilitates greater GPR126 activation than dynamic forces alone and that dynamic force signaling requires the *Stachel* sequence to activate GPR126.

These in vitro findings suggested that laminin-211 enhances GPR126 structural changes to allow for *Stachel*-mediated signaling, and the overexpression studies in zebrafish demonstrated that increased *lama2* levels can drive Gpr126-dependent myelination. What physical forces might laminin-211 impart upon GPR126 in vivo? Interestingly, laminin-211 polymerization is known to be essential for Schwann cell development; *Lama2<sup>dy2j</sup>* mouse mutants have a spontaneous mutation that leads to aberrant splicing and deletion of the laminin  $\alpha 2$  polymerization domain, and these mice have defects in Schwann cell development and myelination [174–176]. To test if laminin  $\alpha 2$  polymerization could be an activating force on Gpr126, reminiscent of physical forces in vitro, Petersen and colleagues engineered a non-polymerizable Lama2 overexpression construct to mimic the *Lama2<sup>dy2j</sup>* mouse mutation. Unlike wild-type *lama2*, the polymerization-defective *lama2 (dy2j)* was not sufficient to rescue myelination defects in hypomorphic *gpr126* zebrafish mutants [155].

In summary, the in vitro studies suggested the possibility that GPR126 might be sensitive to mechanical force, while the in vivo studies implicated laminin-211 polymerization as a potential source to transmit forces and stimulate mechanosensation in vivo.

### 3.2 Latrophilins/ADGRL1–ADGRL3

Contemporaneous work in *Drosophila* has even more strongly implicated an aGPCR in mechanosensation, as latrophilin/CIRL was shown to shape the response of chordotonal sensory neurons by determining the sensitivity of certain mechanosensors in a cell-autonomous manner [59]. Chordotonal neurons are peripheral, compound mechanosensory neurons in *Drosophila*, which perceive mechanical signals such as sound, touch, and muscle stretch [47, 52, 177]. Latrophilin/CIRL is robustly expressed in these cells (unpublished data) suggestive of a function in this cell type. Indeed, analyses of a newly engineered *dCirl* null allele revealed many interesting phenotypes consistent with a role for latrophilin/CIRL in mechanosensation. *dCirl* mutant larvae exhibited aberrant crawling patterns, and although locomotion is a complex behavior controlled by both motor outputs and peripheral sensory inputs, this defect could be partially rescued by chordotonal neuron-specific expression of *dCirl*. Moreover, *dCirl*

mutant larvae were also less responsive to gentle touch (touch perception is a known function of the chordotonal organ), and this phenotype could only be suppressed by chordotonal neuron-specific re-expression of this aGPCR [59]. Interestingly, the morphology of chordotonal neurons is grossly normal in *dCirl* null mutants, suggesting that latrophilin/CIRL is required specifically for the function of these neurons and not for their development or morphology.

To elucidate the potential role of latrophilin/CIRL in mechanosensation, Scholz and colleagues directly applied mechanical stimulation (vibration stimuli) to the cap cells of the chordotonal organ and simultaneously recorded action currents from chordotonal neurons. While *dCirl* mutant neurons maintained spontaneous activity in the absence of vibration stimuli, the mutant neurons displayed significantly lower action current frequencies across the entire tested vibration spectrum. Compound mutant analyses also demonstrated that *dCirl* genetically interacts with TRP channels, known as mechanosensor molecules of *Drosophila* chordotonal neurons [59] (see also Sect. 2.2.3).

All in all, these studies establish that latrophilin/CIRL can modulate the sensitivity of neuronal mechanosensation (Fig. 1). Given that several cell-specific rescue experiments demonstrate that *dCirl* functions cell autonomously in chordotonal neurons, likely via CTF signaling, it will be interesting to determine if and how *Stachel* sequence signaling modulates these functions of latrophilin/CIRL in mechanosensation (Fig. 3).

### 3.3 GPR56/ADGRG1

The role of aGPCRs as mechanosensors also extends beyond the nervous system. Skeletal muscle can respond to multiple stimuli, including mechanical tension, which is a potent regulator of muscle mass. In myotubes, a splice isoform of the peroxisome proliferator-activated receptor gamma coactivator 1-alpha (PGC-1 $\alpha$ ) induces muscle hypertrophy. Interestingly, overexpression of PGC-1 $\alpha$  increases *Gpr56* expression, and GPR56 overexpression also induces muscle hypertrophy dependent on G $\alpha_{12/13}$  signaling. Resistance exercise causes muscle hypertrophy, but this phenotype is attenuated in *Gpr56* mutant mice [156] (see also [178]). These results collectively highlight a potential role for *Gpr56* in sensing muscle fiber size, which could be mediated by mechanical sensitivity to stretch (Fig. 1).

Furthermore, in the central nervous system, GPR56 is required for proper cortical development (i.e., cortical folding), which has been directly linked to mechanosensation [179]. GPR56 and other aGPCRs including GPR114/ADGRG5 and GPR97/ADGRG3 are also highly expressed on lymphocytes [180]; it is tempting to speculate that signaling via these aGPCRs might be affected during the rolling and extravasation behaviors characteristic of these cells.

### 3.4 Mechanosensation by Other Adhesion GPCRs

These recent studies implicating Gpr126, latrophilin/CIRL, and GPR56 as putative mechanosensitive receptors raise the interesting and obvious question as to whether other aGPCRs might share this mechanosensitive property.

Indeed, one splice variant of GPR114 is sensitive to mechanical stimulation [166], although it is not yet clear in what physiological context this receptor may partake in mechanoreception. In addition, recent reports implicate *EMR2/ADGRE2* in vibratory urticaria [181], an autosomal dominant disorder that is signified by hives and systemic manifestations in response to cutaneous vibration. Interestingly, mast cells express EMR2, and mechanosensing through this receptor appears to depend on the stability between the NTF and CTF heterodimer of EMR2. A human mutation in the GAIN domain destabilizes this interaction and conceivably results in increased receptor activity upon mechanical challenge, which ultimately leads to massive mast cell degranulation [181]. Moreover, observations from previous studies on several aGPCRs can be interpreted in the light of these new insights to support a unifying model of mechanical susceptibility of this receptor class.

In the murine lung, loss of *Gpr116/Adgrf5* causes reduced surfactant uptake, leading to massive and pathological accumulations of surfactant lipids and proteins in the alveolar space [182–184] (see also [185]). Given that surfactant is essential to temper lung surface tension to allow for effortless lung expansion during inhalation, it is enticing to speculate that GPR116 is involved in monitoring lung tension (Fig. 1).

Similar to the expression of dCIRL in mechanosensory cells, GPR98/VLGR/ADGRV1 localizes to inner ear hair cells [186], where it is involved in proper development of the cochlear organ of Corti [187]. In the future, it will be interesting to determine the exact role of GPR98 during auditory mechanotransduction in these cells.

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## 4 Conclusions

In sum, ionotropic mechanosensors are distributed throughout various organ systems across different species where they assure mechanosensory responses with latencies on millisecond timescales, rather atypical for metabotropic mechanosensors. Interestingly, however, in some instances these responses appear to be adjusted through the activity of GPCRs, which begs the question if ionotropic and metabotropic mechanosensors form functional units to enable reliable mechanotransduction.

aGPCRs regulate a multitude of physiological processes signified by the capacity to sense mechanics in different cellular contexts, which could explain the participation of aGPCRs in various, seemingly unrelated, biological phenomena. The structural and functional layouts of aGPCRs potentially reflect their optimization for the perception of mechanical forces, rather than a general sensitivity to a range of multiple sensory stimuli known for other GPCR families. Another

interesting aspect will be to unveil the signaling cascades through which aGPCRs influence ionotropic mechanotransduction processes and whether ion channels and GPCRs form functional mechanosensory units. Further research efforts will clarify the physiological and pharmacological properties underlying the mechanobiological roles of aGPCRs.

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