



TGF- β Family Signaling in Ductal Differentiation and Branching Morphogenesis

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Epithelial cells contribute to the development of various vital organs by generating tubular and/or glandular architectures. The fully developed forms of ductal organs depend on processes of branching morphogenesis, whereby frequency, total number, and complexity of the branching tissue define the final architecture in the organ. Some ductal tissues, like the mammary gland during pregnancy and lactation, disintegrate and regenerate through periodic cycles. Differentiation of branched epithelia is driven by antagonistic actions of parallel growth factor systems that mediate epithelial–mesenchymal communication. Transforming growth factor- β (TGF- β) family members and their extracellular antagonists are prominently involved in both normal and disease-associated (e.g., malignant or fibrotic) ductal tissue patterning. Here, we discuss collective knowledge that permeates the roles of TGF- β family members in the control of the ductal tissues in the vertebrate body.

TGF- β FAMILY MEMBERS IN DUCTAL MORPHOGENESIS AND EPITHELIAL–MESENCHYMAL INTERACTIONS

The epithelial cells contribute to the development of various vital organs by generating tubular and/or glandular architectures. The fully developed forms of ductal organs depend on processes of branching morphogenesis, whereby frequency, total number, and complexity of the branching tissue define the final architecture in the organ. Some ductal tissues, like the mammary gland during pregnancy and lactation, disintegrate and regenerate through periodic cycles. Differentiation of branched epithelia is driven by antagonistic actions of parallel growth factor systems that mediate epithelial–mesenchymal communication. Transforming growth factor- β (TGF- β) family members and their extracellular antagonists are prominently involved in both normal and disease-associated (e.g., malignant or fibrotic) ductal tissue patterning. Here, we discuss collective knowledge that permeates the roles of TGF- β family members in the control of the ductal tissues in the vertebrate body.

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body (2001). Although the role of the TGF- β family in the development of the digestive tract is well established, the role of the TGF- β family in the development of the reproductive tract is less clear. In the mouse, the TGF- β family is expressed in the developing reproductive tract, and the TGF- β family is involved in the development of the reproductive tract (McNally et al., 2013). In the mouse, the TGF- β family is expressed in the developing reproductive tract, and the TGF- β family is involved in the development of the reproductive tract (McNally et al., 2013). In the mouse, the TGF- β family is expressed in the developing reproductive tract, and the TGF- β family is involved in the development of the reproductive tract (McNally et al., 2013).

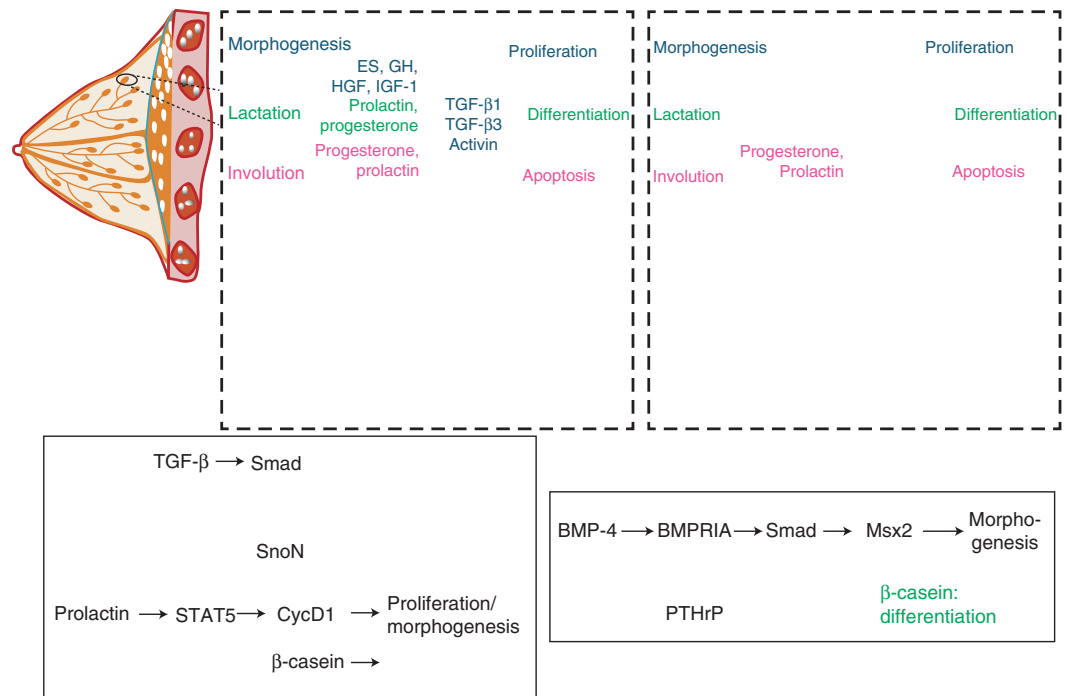
Adding to the complexity of the TGF- β family, the TGF- β family is involved in the development of the reproductive tract. In the mouse, the TGF- β family is expressed in the developing reproductive tract, and the TGF- β family is involved in the development of the reproductive tract (McNally et al., 2013). In the mouse, the TGF- β family is expressed in the developing reproductive tract, and the TGF- β family is involved in the development of the reproductive tract (McNally et al., 2013). In the mouse, the TGF- β family is expressed in the developing reproductive tract, and the TGF- β family is involved in the development of the reproductive tract (McNally et al., 2013).

THE TGF- β FAMILY IN GLANDULAR ORGAN DEVELOPMENT AND EMT

Mammary Gland

The development of the mammary gland is a complex process involving the TGF- β family. In the mouse, the TGF- β family is expressed in the developing mammary gland, and the TGF- β family is involved in the development of the mammary gland (McNally et al., 2013). In the mouse, the TGF- β family is expressed in the developing mammary gland, and the TGF- β family is involved in the development of the mammary gland (McNally et al., 2013).

Adding to the complexity of the TGF- β family, the TGF- β family is involved in the development of the mammary gland. In the mouse, the TGF- β family is expressed in the developing mammary gland, and the TGF- β family is involved in the development of the mammary gland (McNally et al., 2013). In the mouse, the TGF- β family is expressed in the developing mammary gland, and the TGF- β family is involved in the development of the mammary gland (McNally et al., 2013). In the mouse, the TGF- β family is expressed in the developing mammary gland, and the TGF- β family is involved in the development of the mammary gland (McNally et al., 2013).



ab f, e c a c e e c e e d
 BMP-4 e I a e d b e a a d
 e e e a e d e (PTHrP), i c d i c e
 e e f, e BMP e I e c e BMPRIA
 b c e e c a c e (F.1). I e a b e c e f
 BMP-4, e c a c a b d f a d e e f

a d e b a c c e e e e
 I b e d (He e a. 2007). BMP a c e
 c a c a d i c a c e e a a e a e,
 a a a e b I d e f, e e a c e I a BMP
 a d a c c e e, i e d a I a
 (Tü) (F c a e a. 2013). Tü e e e d

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 a ced a d fa e b ea ca ce \rightarrow e a
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 \rightarrow a e e a ce e a e bee d e d f \rightarrow
 \rightarrow ce e e a d b c \rightarrow e \rightarrow \rightarrow
 \rightarrow I e a e e e a e e a I e e e
 \rightarrow I a \rightarrow a I -40 a e Ta e \rightarrow ce
 ac \rightarrow e c S ad e e (K e a. 2010).
 I e e e \rightarrow a e e a ce , TGF- β
 d ce ce -c c e a e , a a d EMT
 S ad3, b I S ad2, e e a a
 e a d \rightarrow a e e e I e b
 S ad2 a d S ad3 (K e a. 2010). T e e
 d I a e e \rightarrow a ce f e I a
 TGF- β a d e a c a d a c a
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 a a d e \rightarrow a a d d f f e e a
 a d a c a , I c a ac a b e
 Ja I a e (JAK) a d e a a d ce
 a d ac a f a c 5 (STAT5) a
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 (d e f e ac β A c a , a i a
 e b - β A c a) (Nac i a e a d B. i
 2016) c a ac a e S ad2, S ad3, a d S ad4
 a c i e e , e f e e i
 e a c a f STAT5 a d c ac a
 CREB-b d e (CBP) 300, a
 S ad e f \rightarrow e b c \rightarrow e
 e a e c ac a (C c a e a.
 2008). I \rightarrow a e , ac -a d TGF- β -d
 e S ad c e e c e f a c a
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 a c f STAT5 a e e e , e d
 e e e c d β -ca e a d c c DI (F .
 1). A e c d e a \rightarrow e a d
 d f f e e a ce \rightarrow e d a d b e S ad

cellular S. N (Jagoe et al. 2012). S. N
 e e e e a c a ac f S ad
 c e e, a d TGF- β a f d i ce e
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 (Jagoe et al. 2012). I a e e e i
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 a c fac f e l c e a S ad c e
 e c c a a c a e d e a e c
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 TIF1 γ), f e l a e l b e f TGF- β -e
 e e e (Ma a l e 2012). Ma c a e
 e a c e c a b a f e T i 33 e e
 e l a c a c a a d d e e c e
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 c d e, TRIM33 b e S ad a d STAT5
 a a c a e f a a d e e e e e
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 a e a c e (He e a. 2013). T l e
 e a e e f f e c f TGF- β d i e a e a e
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 a c STAT5 a i a, i e e a e e a l c e a
 c e l a a e a e a a c c
 a b e i e e e e l a i a
 I c a c a c a c e, EMT, a b e e i
 a d c e l a a a e d l c e. I l d e
 (e e i e d M l a a a d H e d 2013). R e
 c e l a e l a f e EMT c e, l c
 a a c fac a d e c c c RNA
 (c RNA) e c a b e a a e d d e e d e
 a e c f e c a c b l
 l a c a a d d e e l e l d e e
 c e f c a c e e. Ma c a EMT
 e l a e d b c c e c e a c l e. F
 e a c e, c e l e f c a c a e e
 a c e TGF- β b a e e c a c a d
 a c e e f e e c e, a d c a
 a a e c c e c c a (EMT) e
 e l e a c e d f e a l e f c e a (Ga
 e a. 2008). I add , e d f e e

ace l a c c e c e e
 a l l f TGF- β , a a i d
 f f e a i c a c a d a f c e
 e, i e e a f f e c e e
 EMT e e (Le e e a. 2012). T e a
 c fac e c e 2 (OVOL2)
 e e a f e c a c a d i c a c
 e e a d c a e a c e f a d e c e
 e c a c a a d (Wa a a b e e a. 2014).
 OVOL2 e e e e f c a EMT
 a c fac a d l e e a f
 TGF- β d i c e c a c a e e a c e EMT
 (Wa a a b e e a. 2014). De e OVOL2 f c
 e c a c a e e l c b c a c a
 d i c d e e c e, c a l e f f e c e
 a d e a c e TGF- β -d e e d e EMT, a e
 l b c a d f f e e a f e a d (Wa
 a a b e e a. 2014). T e c e f MET f e
 d e b BMP a d a b e e e d e
 e e c a c a e e a c e f c EMT, l
 c b l c a c a e e a d f f e e
 a (M l a a a d H e d 2013).

Prostate Gland

A d e c c e d e d e e c e f
 e c e a (T c a d Ma e
 2006). T e e f a c d f f e a
 d e d a a c e a b e e
 e e a a d a e d l e e c l a c e
 l c TGF- β fac c e b e l c a e
 c e e e (T c a d Ma e 2006).
 T e e e TGF- β , a c A, BMP-4, a d BMP-
 7, b a c d e e c e, i e e a
 l a d d f f e e a fac (GDF)-7
 c e l a d b a c c e e
 (T c a d Ma e 2006).

TGF- β 1 c a e a e e a c e e c
 c a c a c e a c a d f c l e
 e c b a d e l b (T c e e a. 1996).
 Q e e e f TGF- β 1 a c e e
 a c e e l a e a c e e e a
 e e c b e c a c e e e e a d
 (T c e e a. 1996). O e e e a d, TGF-
 β b c a e a d d f f e e a b
 c c e c c e a e l l d i c f
 c c -d e e d e a e b 21 e
 e, a d e l b a f f c e b
 e e e a a f e b e l d e a e d



c < e . f . a e d e e < e . a
 (Ç a e a . 1999). S i c a a b e e f f e c t f e
 TGF- β , e . e e e . e a . a e c
 e - d e e d e . T e i i b e f f e c
 < a . e < < a I e e e c e
 d e e . b i d , i . e e a TGF- β < a e e
 < I a e . f e a . f d f f e e a e d e e
 a (T < e a . 2004); i . c < a e a
 i . I c e c a c . (T < e e a . 1996).
 T e a e i < . e e c a d e f
 TGF- β . a . e . < a e i . a e
 c e d e c e e d i a i i
 b . e . e (S a e a . 2005). T e a c
 i TGF- β . a c e c e a d
 e . i . f e a . a e a . < a a e d b
 N c l a . a e d e . e e c e e
 e TGF- β b d i c e e . f e
 TGF- β e I e c e i (T B R I) (V a d e e a .
 2012). T e TGF- β a d e < a a e d b
 e I a . a c . f a d . e , a d c a e
 a d . e a b i d a c e c a e e i c . e a d
 d e a . e d e c . (S a e a . 2005).
 I a e e e i . e e c . a . e f e
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 e a d f f e e a . a d c . e e , c i
 c i e . f c . I e b a d d e I . e a I e i
 < e e c < a c e . f e . a c i . d i c
 (I . e a I) , d i c e . e e e a . f
 a c e . e a c e (L e a . 2009). H i e
 e , b a d d e c e i a c T B R I I . I e
 e c c e e a c . a . e c a . e a a d
 c a . a d f f e e i a e (L e a . 2009). T e
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 a e e . e a W a . e a d . e
 e e a . f . a c e . e a e c e (L
 e a . 2009).

S c a , e . c i I e f a . a c
 e . a i . a c . A . i a . b . c
 b a c . i . e e . f e d e e
 d i c , i . e e a . e a a . f i a
 < e b a c . f e d i c (C a c i a e a .
 2001). N c a d e e . a . a e a d
 e . e a c . - β_A (a c . - β_B a e a < a
 d i I b e) , f a . a d . e a c . i e
 I a d e I I e c e i (C a c a e a . 2001).

I a d d . TGF- β , BMP-4
 e e d a e . e a < e e c < a b d e
 e d e e . a c a d . BMP-4 < a
 c . b a c . < . e e , b e c a I e

< c e i . e B 4 e e c . i e
 a c e d b a c . (L a c e a . 2001). C i I e
 f . a e e . e a c e . e e e c e f
 BMP-4 . i . b . f . f e a . a d a
 b . c . b i d d i c a c . (L a c e a . 2001).
 BMP-7 < a a c < a . a BMP-4 , a d . a e
 a d f . < B i - / - < c e a e e a e d i
 e e . e b a c . (G . a e a . 2005). I e
 . a e . a c i I e BMP-7 d i c e e
 e . f b a c . b i d , i . c < a . e
 c a e c . f N c l a a c e i a
 (G . a e a . 2005). T e a c . f BMP
 a d i c a b e c e d d i . a c a d d e
 e . < e b . e d e a c e a e . a e e
 e e e . f . e B 2 a d B 4 e e
 (K e e a . 2015). A . e e e . e a . e e
 a c e I a a c . f BMP . e d e
 . a e a e b a a c e d b a . a . , I c i a
 . I N g - / - < c e , a c . i . e
 e . , e . e a . e . f . e . a c
 a d e e e b . e . a . c a c . f
 e BMP . a . a e c e . e . e a b e c e
 f . e e a . e e I a . e e e d b
 (Ç . e a . 2007). BMP . a . e . a e
 e . e I c . c e d a e d b BMPRIA . C d
 . a d e e . f . e c e . e I . e a
 I e . e a c e . b . c c e d f f e e a .
 a d . e . a c c e . e e . e . f . e
 a c . f a c . N 3.1 , i . c < a
 < e . a c e . e a d f f e e a . b
 BMP (O c . e a . 2014).

I c . a . e . e . e < e c b e f . e
 f a c , GDF-7 (a . i . a CDMP-3
 BMP-12) < e . a e e . e a d f f e e
 a . (S e e e a . 2001). E i . e c i I a
 a a e . f . e c e a < . f a c . f GDF-7
 . e . a e e a . b e e f c e d .

Pancreatic Organogenesis

T e a c e a . a e d d e c - d e e d . a
 . a c . f . e e c a c e e a e . e e
 c e , d i c a , a d e d c e c e (K c a d H e b
 . 2001). T e e c e . e a e . a e f c
 . e a c e a c b i d . a . a . I . i . f c
 e f e I , < a . e . e b i d (e e b e i)
 (K c a d H e b . 2001). T e < . d a a
 c e a c c e I d e . c c e . f EMT a d MET
 d i . f e a . a d c . e e .

... a ... d a e ... e a c e ... e e a e < e e -
 < a' c e ... a e' e ... a ... e e a e d f f e e -
 a e d e ... e a a d e c e ... c e (T ...
 a d M a' e' 2006). TGF- β ... b ... e e a
 d f f e e a ... e a c e a' e' < ... e' e
 ' a d I a' e' a' . E ... e a ... e e' e ... f a
 d < a - e a ... e < I a' T β R II ... a' e c
 < c e' e I ... e ... f e a ... a' d a b ... < a
 d f f e e a' ... f a c' e a c' a c (B ... e
 e' a. 1997). T ... I , TGF- β ... b' e' c' e' a -
 c e a c c e d f f e e a' , I ... e e a e' a c e d
 TGF- β 1 e ... e' e a d ... a ... b ... a d
 e c e ... e a ... e e (B ... e e a. 1997).
 E ... c I I e' f < I e e x b ... c a c e a c
 ... e a ... e ... f ... e d f f e e a ... f
 ... e a ... I c e ... e a d ... e a' c ... e c I a ...
 a a' ... e d e e' ... a ... d . U -
 ... < e' d ... , TGF- β 1 i a ... i
 ... b' d f f e e a ... f ... e a c a < c a -
 < e' f ... e' a' d (S a ... e' a. 1994). A -
 ... I ... TGF- β 1 a c' a a ... b ... f d f f e e -
 a ... f e ... c e a c e a c' e' ... e a' c e ,
 ... e e d c' e β - ... e' c e' d f f e e a' ...
 ... e ... a' d c I I e d e' ... (S a' ...
 e' a. 1994), ... c' < a b e i' ... e a c' ...
 f TGF- β d I' ... a c e a c d e e ... < e' .
 H ... e e , e ... e' ... f ... e d' < a - e a ... e
 T β R II ... e a c' e a c e ... e I' I ... e' a
 TGF- β ... a ... b' e' d' c' e' a c e a c d
 d f f e e' a' ... ' ... (T i a' a' e a. 2007). A
 < a ... b' ... b' TGF- β ... a ... a
 a' a e' d' ... a ... f ... e e' d' c' e' a c e -
 a c' a d (D ... a' e a. 2016). B' a c' a
 S' a d 3, TGF- β ... < e ... e e ... e' f' e
 c e - c c e e I a ... 16¹ 4a, ... c' c' ... e
 e f - e e i a ... f ... e a d I ... e d' c' e' e x c e
 ... e a c e a (D ... a' e a. 2016). U d e' e
 a c' ... f TGF- β , 16¹ 4a a c c I < I a e , ... b -
 ... I e e e e a ... a ... e a < a' a e
 (D ... a' e a. 2016). B a e d ... < e' c' a < ,
 ... a < a c ... c a ... b ... f ... e T β R I ... a e
 ... e < a ... a' a' ... f e d c' e β -c e' ... a
 c a ... e' e a' e ... d e a < a' (D ... a' e a.
 2016).

TGF- β 1 ... b ... d I c a e ... e a < ... -
 e e ... < a' a' a c e a c' e' e' a ... c I -
 I e' e' x' b e d d e d ... e e - d' < e' ... a c' a e
 < a c e ; ... e e' , a - TGF- β a' b' d ...
 < e' d I c' a < ... e e' , b' e' I a ...

TGF- β e c e e d b ... e a c e a c' I d < e
 (H a ... e a d R ... e b e 2007). S' < I a' ... f
 a c e a c d I c' c e ... a e d f' < ... I < a
 d ... I ... TGF- β 1 ... d I c e E M T a d' a -
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 e' c' c' a c ... f a c ... , I ... c' < a -
 b e' ... a ... b' ... e f TGF- β ... e d -
 c' e' c e d f f e e' a' ... a d a b ... < -
 < e' a d f f e e' a' ... (S' ... e' a' . 2011). I
 a e e x e' i' ... e' a' e' b' e' a' ... , I < a
 e ... e a' c e a c c e I d e' E M T i' e
 c I I' e d ... (L ... a e a. 2013). T ... e e x e -
 e' c' < a' d e' a' e' c a b e ... a x c e d
 ... a d a - e' c e' f' ... a d e ... a e' e -
 ... f e d c' e a c e a c' a c' ... f a c -
 ... (S' ... e' a' . 2011). I ... c' a , β - ... e' c e
 c a b e' e e a e d ... I ... a c' ... a e ...
 a x < ... , b' I ... a R ... - a' c' a e d ... a e
 a d a TGF- β e c e ... a e ... b' ... (L ... a
 e a. 2013). T ... I , TGF- β - d I c e d E M T f' a -
 c e a c e ... e a c e < a' e' e e a e , I < a
 a c' e a c' c e' ... a c a ... e b e a' a e d ...
 d a b e c' a e ...

S' ... a' TGF- β , a c ... A b ... c' b a c' -
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 ... e f ... a' b' ... c' a c' ... , a d ... I ... -
 < ... e' a' c' e a c' a c' a' c' e' d f f e e' a' ...
 ... d e c e a e d e' d' c' e' β -c e' ... < b e' (R -
 ... e a. 1995; Z ... a' e a. 2004). H ... e' e ,
 ... e , e < a , e ... e < e' ... a c e a c
 e ... a' ... I' e d ... a a c' ... A' a c' ... B (a c -
 ... - β B d < e) d d ... f a f f e c' a' c' e a c c e
 d f f e e' a' ; ... e a d , a c' ... A c a' e d a
 ' a d f f e e' a' ... f ... e e x' b' ... c a c e a -
 c' ... e' ... e' a' I e (a E' e a. 2004).
 I' d e' e' d e' ... a' d f f e e' a' ... a d' a
 e ... e < e' c' ... < e d , a' a c' ... A' a c' ...
 B' I' e' e' e' c' e' α -c e' d f f e e' a' ... a' d
 ... e e d c' e' β -c e' d f f e e' a' , e -
 a b ... I' e' c e' ... (A' d e e i' ... e a.
 2015). T' e' a c' ... e' e' e' e' e' e' ... f
 α -c e' ... a c' ... f a c' ... a d' d I c e' β -c e
 ... a c' ... f a c' ... e' e' ... , a d' e' a c -
 ... e' ... a' e' e' ... e' f' < c e i' ... a d e -
 e' ... f' ... e' F I 3 e e , e c' d' I' ... a' - e
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Salivary Gland Formation

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[illegible]

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THE TGF- β FAMILY IN GASTROINTESTINAL TRACT DEVELOPMENT AND EMT

Esophageal, Stomach, and Intestinal Differentiation

Along with BMP, TGF- β is a key signaling molecule in the development of the gastrointestinal tract. TGF- β is expressed in the developing gut and plays a critical role in the differentiation of the esophagus, stomach, and intestine. TGF- β is secreted as an inactive complex with its binding partner, TGF- β RII, and is activated by proteolytic cleavage. The active TGF- β complex then binds to the TGF- β RI and TGF- β RII, leading to the activation of the TGF- β signaling pathway. This pathway involves the recruitment of Smad proteins, which then translocate to the nucleus to regulate gene expression. TGF- β is also involved in the regulation of cell proliferation and differentiation, and has been implicated in the development of various types of cancer.

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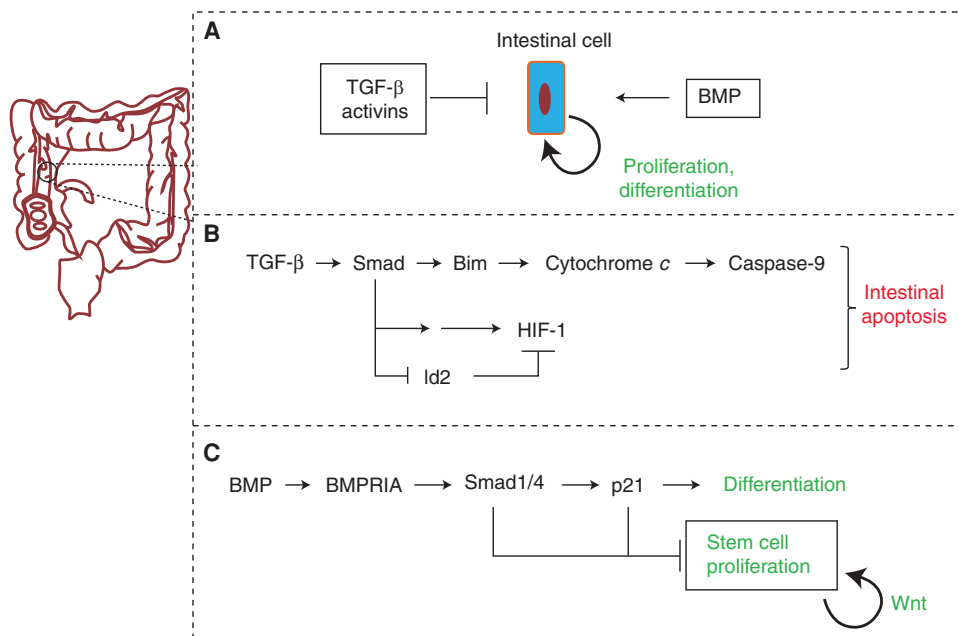


Figure 2. Regulation of TGF- β family signaling in the gastrointestinal tract. (A) TGF- β and BMP signaling in an intestinal cell. (B) TGF- β signaling pathway leading to intestinal apoptosis. (C) BMP signaling pathway leading to differentiation and stem cell proliferation.

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THE TGF- β FAMILY IN LIVER AND BILE DUCT ORGANOGENESIS AND EMT

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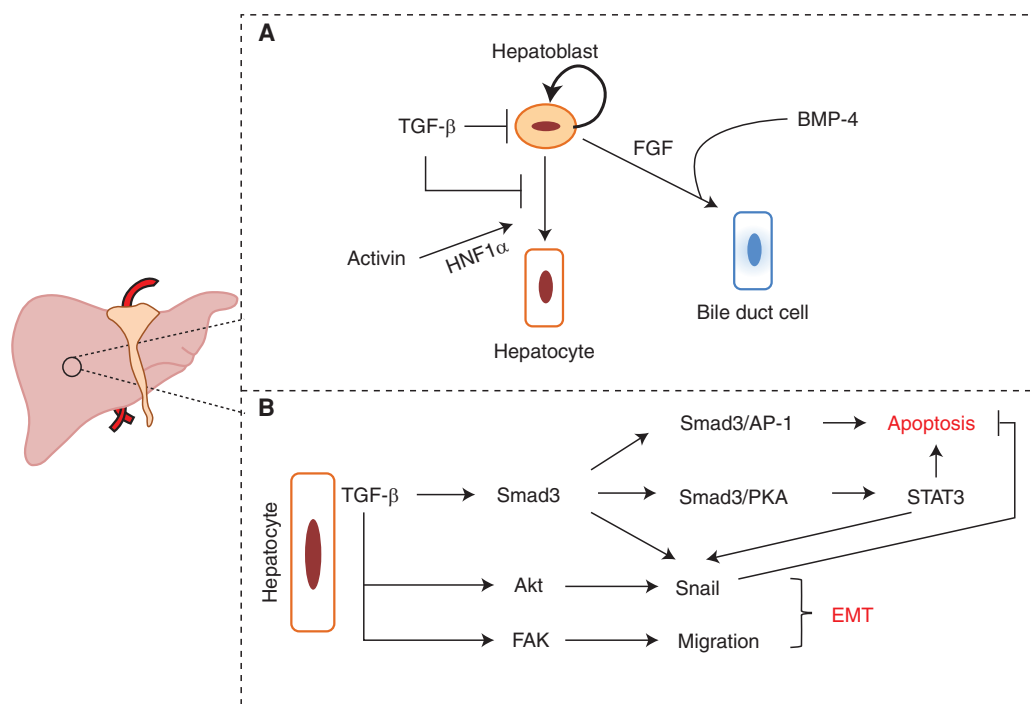


Figure 3. Regulation of TGF- β signaling in liver development and EMT. (A) TGF- β signaling promotes hepatocyte differentiation and inhibits hepatoblast proliferation. Activin and HNF1 α promote hepatocyte differentiation. BMP-4 and FGF promote bile duct cell formation. (B) In the liver, TGF- β signaling activates Smad3, which then interacts with AP-1, PKA, and STAT3. Akt and FAK also contribute to the EMT process, leading to Snail-mediated migration and EMT. Apoptosis is also regulated by Smad3/AP-1 and STAT3.

[illegible]

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 a c e d a e d b GPI-a c e d c e c e
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f' e BMP-4 e e a d e c e . b' < e-
d e < a c e , I . < d e e . c' < e d e <
< . e e a' d I b' e I e d' f f e e a .
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c a e d b e a d . e x b . c e' a c a f
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I e , < < . a BMP a e c a f . e
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d e < , . a e . c h I d < . e e c a
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b . I' c' d e e a' , BMP-2 a d , I e e c-
e d , BMP-4 b I b I a . 1 . a d b a c -
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e a. 1997; R a a' a e -A . a e a' . 2000).

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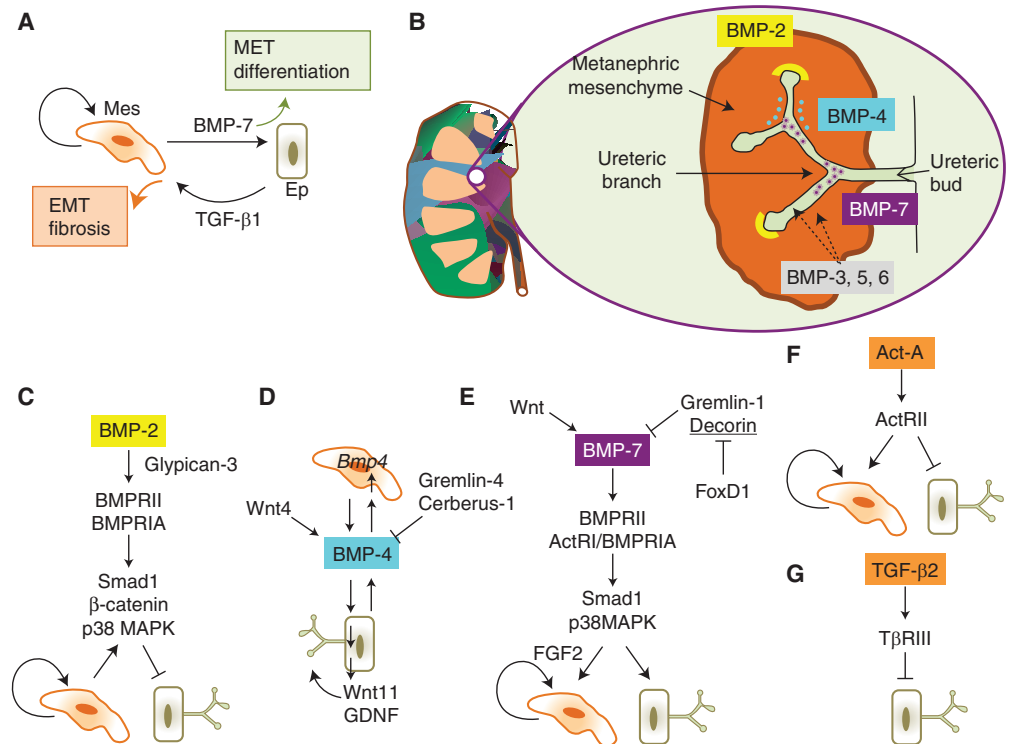


Figure 4. Regulation of TGF-β factors in kidney development. (A) Schematic of EMT and MET differentiation. (B) Kidney development with BMP signaling. (C) BMP-2 signaling pathway. (D) BMP-4 signaling pathway. (E) BMP-7 signaling pathway. (F) Act-A signaling pathway. (G) TGF-β2 signaling pathway.

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K. Kahata et al.



ced BMP (Hüe et al., 2009). Addition of BMP to the culture medium induces BMP signaling, leading to the expression of BMP target genes (Hüe et al., 2009).

The BMPRIA gene encodes a type I receptor, which is a transmembrane protein with an extracellular domain, a single-pass transmembrane domain, and an intracellular domain. The intracellular domain contains a kinase domain, which is responsible for the signaling activity of the receptor. The BMPRIA gene is expressed in a variety of tissues, including the brain, heart, and muscle. The BMPRIA gene is also involved in the regulation of bone development and the differentiation of muscle cells (Ebner et al., 2006). In addition, the BMPRIA gene is involved in the regulation of the BMP signaling pathway, which is a key component of the BMP signaling pathway (Ebner et al., 2006).

BMP-4 is a member of the BMP family of growth factors. It is a dimeric protein composed of two identical subunits, each of which is a single-chain polypeptide. BMP-4 is secreted by a variety of cells, including fibroblasts, epithelial cells, and endothelial cells. It binds to the BMPRIA and BMPRII receptors, which are type I and type II receptors, respectively. The binding of BMP-4 to these receptors leads to the activation of the BMP signaling pathway, which is a key component of the BMP signaling pathway (Ebner et al., 2006). BMP-4 is involved in a variety of biological processes, including bone development, muscle differentiation, and the regulation of the BMP signaling pathway (Ebner et al., 2006).

Receptor-mediated BMP signaling is a key component of the BMP signaling pathway. It involves the binding of BMP ligands to BMP receptors, which leads to the activation of the BMP signaling pathway. The BMP signaling pathway is a key component of the BMP signaling pathway, which is involved in a variety of biological processes, including bone development, muscle differentiation, and the regulation of the BMP signaling pathway (Ebner et al., 2006).

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