

**Osteocytic Wnt-TGF β signaling transduction axis activates RANKL
transcription for osteoclastogenesis**

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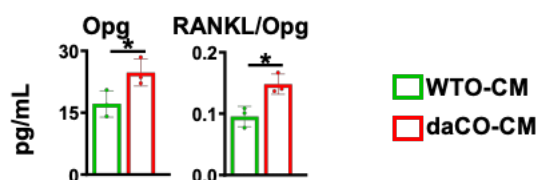
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Supplementary information

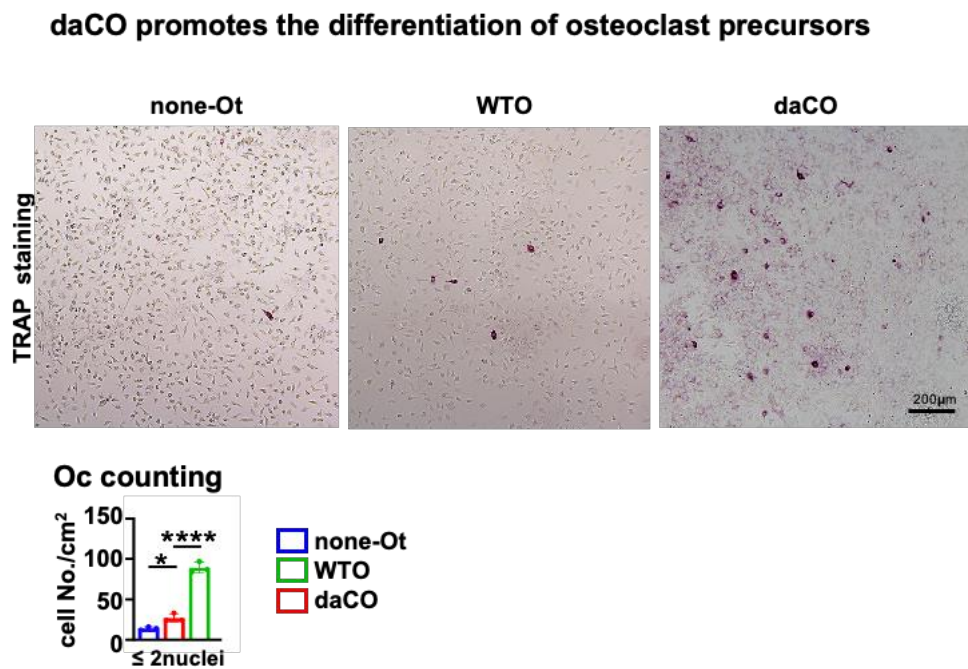
Supplementary information, Figure. S1

detection of Opg and RANKL/Opg ratio in daCO CM



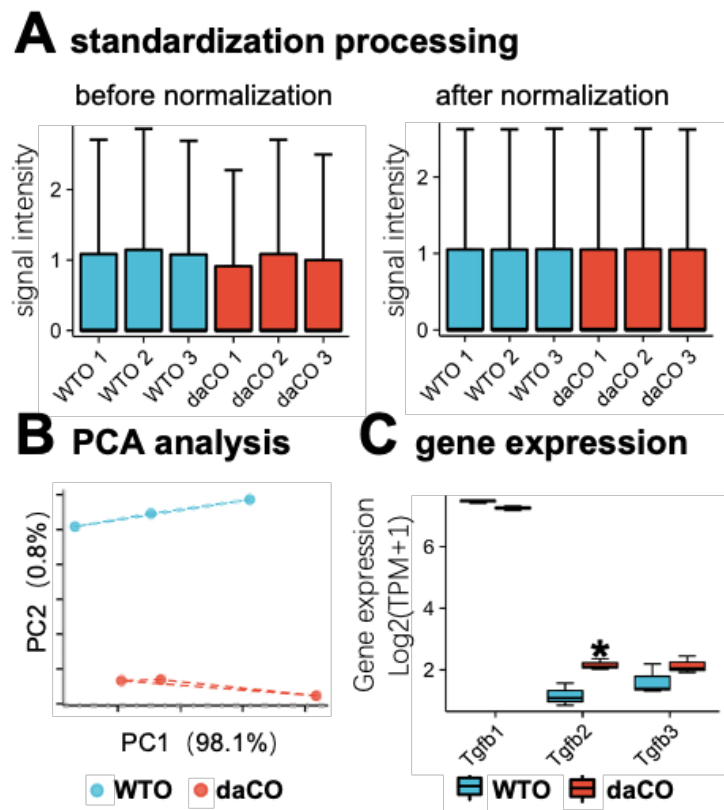
Supplementary information, Fig. S1 daCO conditioned medium has higher concentration of Opg protein with higher RANKL/Opg ratio. Opg protein concentration in osteocyte culture medium as detected by ELISA. Data were expressed as mean $\pm$ SD. WTO, wild-type osteocytes; daCO, osteocytes with dominantly active β -catenin. * $p < 0.05$ v.s. WTO-CM by t -test, $n=3$.

Supplementary information, Figure. S2



Supplementary information, Fig. S2 daCO induces the differentiation of osteoclast precursors in growth medium. TRAP staining of the co-culture of BMMs and osteocytes in growth medium for 11 days and TRAP-positive cell counting. Data were expressed as mean $\pm$ SD. * $p < 0.05$ v.s. only BMMs without osteocytes, # $p < 0.05$ v.s. WTO by One-Way ANOVA, $n=3$.

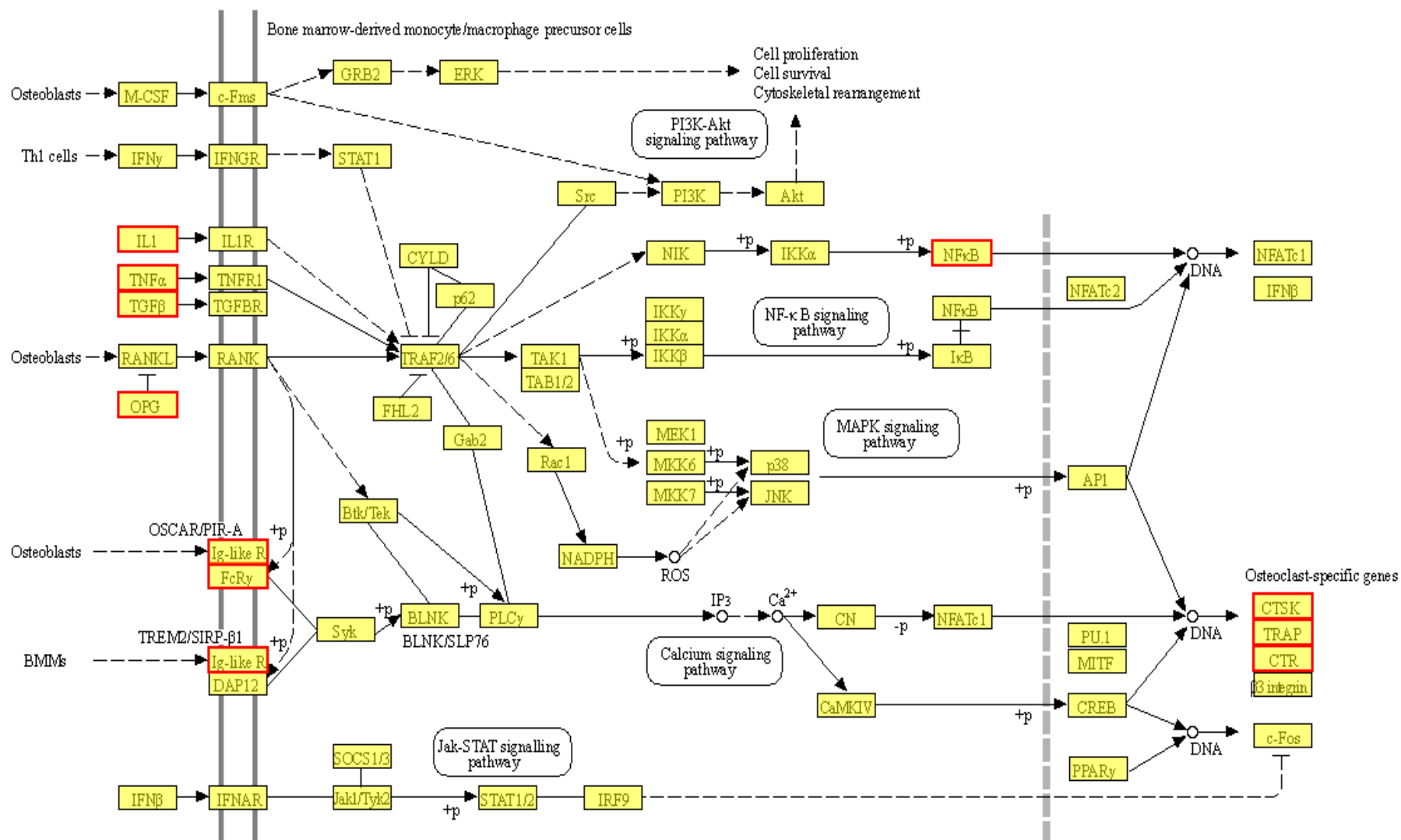
Supplementary information, Figure. S3



Supplementary information, Figure. S3 Samples of WTO and daCO for RNA-seq assay. **A** The baseline levels of raw detected signal before and after standardized using log2+1 normalization. **B** Analyze significant differences and separation between the daCO and WTO groups using PCA dimensionality reduction. **C** The expression of *Tgfb1*, *Tgfb2*, and *Tgfb3* using grouped box plots. PCA, Principal Component Analysis.

Supplementary information, Figure. S4

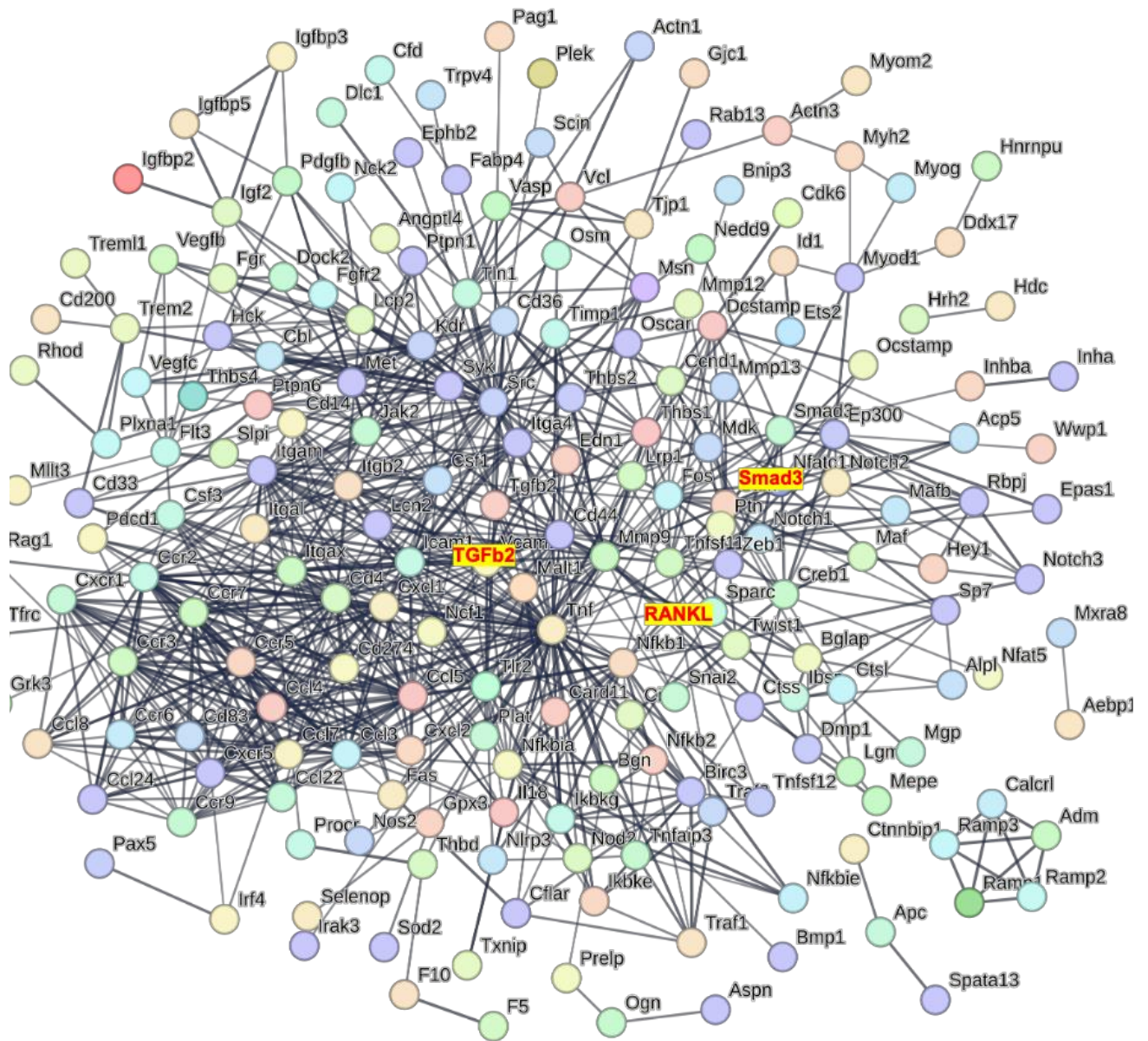
OSTEOCLAST DIFFERENTIATION



Supplementary information, Figure. S4 The KEGG pathways of osteoclast differentiation involves the differentiated expressed genes enriched from daCO transcriptome analysis that are most correlated with osteocytic Wnt. Osteoclast differentiation includes IL-1, TNF α , TGF β , Opg, Ig-like R/FcRy pathways in the upstream of osteoclast differentiation and its downstream, such as NF κ B, CTSK, TRAP, calcitonin receptor.

Supplementary information, Figure. S5

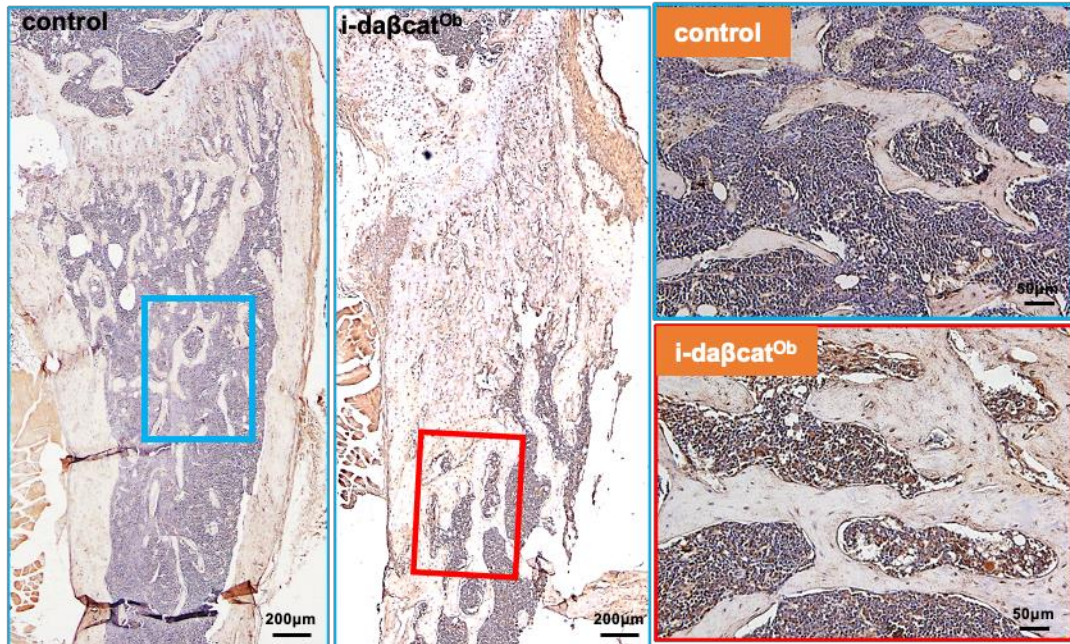
PPI analysis for the proteins related to the differentially expressed OcD genes in daCO



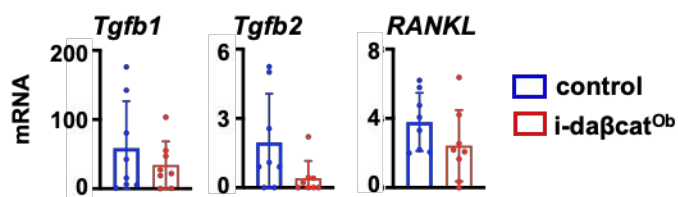
Supplementary information, Figure. S5 Protein-Protein Interaction (PPI) network of the involved differentially regulated genes regarding OcD.

Supplementary information, Figure. S6

A IHC of p-Smad2/3 on mouse femoral section of i-da β cat^{Ob} mice



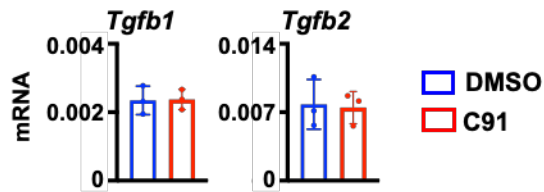
B gene expression in the long bones of i-da β cat^{Ob} mice



Supplementary information, Figure. S6 The effect of osteoblastic Wnt on TGF β signaling. A *Tgfb1*, *Tgfb2*, and *RANKL* expression of the femurs of i-da β cat^{Ob} mice. **B** Immunohistochemical analysis of the p-Smad2/3 of the TGF- β signaling pathway in the longitudinal cross sections of mouse femurs. * $p < 0.05$ v.s. control mice by *t*-test, $n=8$. i-da β cat^{Ob}, tamoxifen-induced expression of dominantly active β -catenin in osteoblasts.

Supplementary information, Figure. S7

detection of TGFb1/2 in primary calvaria cells treated with C91



Supplementary information, Figure. S7 The effect of osteoblastic Wnt on TGF β signaling. Tgfb1 and Tgfb2 expression of the mouse primary calvaria cells treated with Wnt agonist C91. n=3.

Supplementary information, Table. S1

Table S1. Sequences of primers for RT-PCR (mouse)

Primer	Forward	Reverse
<i>Lef1</i>	TACCCCAGCCAGTGTCAACA	TCCATGATAGGCTTTGATGACTTTC
<i>Axin2</i>	TGCAGGAGGCGGTACAGTTC	GCTGGAAGTGGTAAAGCAGCTT
<i>Bmp4</i>	GAGGAGTTTCCATCACGAAGA	GCTCTGCCGAGGAGATCA
<i>Smad6</i>	AAGATGCTGAAGCCGTTGGT	CGAACTCCAGTATCTCCGCTTT
<i>Alpl</i>	ACACCAATGTAGCCAAGAATGTCA	GATTCGGGCAGCGGTTACT
<i>Colla1</i>	GCTGGCAAGAATGGCGAC	AAGCCACGATGACCCTTTATG
<i>Sost</i>	CGTGCCCTCATCTGCCTACTTGTG	CCGGTTCATGGTCTGGTTGTTCTC
<i>Dmp-1</i>	CATTCTCCTTGTGTTCCTTTGG	TCAGTATTGTGGTATCTGGCAACT
<i>RANKL</i>	CATGACGTTAAGCAACGG	AGGGAAGGGTTGGACA
<i>Opg</i>	ACGGACAGCTGGCACACCAG	CTCACACACTCGGTTGTGGG
<i>Rank</i>	TGAGCCTCCGAGCAGAACTGAC	CTGCCTGTGTAGCCATCTGTTGAG
<i>MCSF</i>	CGCTGCCCRTCTTCGACAT	TCTGACACCTCCTTGGCAATACT
<i>Ctsk</i>	GGCAGGGTCCCAGACTCCATC	TGAAAGCCCAACAGGAACCACAC
<i>Dcstamp</i>	TCCTCCATGAACAAACAGTTCCAA	AGACGTGGTTTtaggaATGCAGCTC
<i>Nfatc1</i>	GAGACAGACATCGGGAGGAAGA	GTGGGATGTGAACTCGGAAGA
<i>CHoB</i>	CAGAATGGTAGGAAGGTCACG	CGAATGCTGTAATGGCGTATC

Supplementary information, Table. S2

Table S2. si-RNA sequence

Gene name	Sequence
RANKL-siRNA1	GGATGAAACAAGCCTTCA
RANKL-siRNA2	CAGACTATCTTCAGCTGAT
RANKL-siRNA3	CATGACGTTAAGCAACGGA

Supplementary information, Table. S3

Table S3. ChIP-qPCR primer sequences designed for 5 peaks

Primer	Forward	Reverse
P1	GAGGAGACCACCATCAAGAATCGT	TTGTCTAGTTCTTCCTGGCTCACC
P2-1	AGGTGAGGTGCTGTCCCTTAGAA	CCTGTGGAAGTGTCTGTTAGAGAACT
P2-2	GTCTTGAATCAGAATTGTGCCTTTAGG	GGGACAGCACCTCACCTTTGAA
P2-3	CCTCAATGCTGGGATTAAAGGTATGT	GTCACCTAAAGGCACAATTCTGATTC
P3-1	CGTTTCTCCCATCCAAGTGCTAACC	GCCTCTATCCTGGTGGTGATCTGAT
P3-2	GAACCATCAGATCACCACCAGGATAG	ACAGAGTTGAGTACATACTGACCTTGG
P4-1	AGGCGTCCAACCTATGTAGACTGAAC	ACAGGAGCAGAACCGATAGAATGAA
P4-2	CCACCTTACACTGCTCGGTTTCATT	TCAGTCTACATAGTTGGACGCCTCA
P4-3	TTCATTCTATCGGTTCTGCTCCTGT	GCCAAGAGTTCTGCCTGAGTCTC
P5-1	CTCTAATCCACAGCCTTCCTGACTG	CCTGCTTCCTGACCTGCTTGAG
P5-2	TCAAGCAGGTCAGGAAGCAGGAG	GCTGATGACACAAAGGTGAGGAAGT