



# clone 2.72F, #19

95491™

## Description

Detects:

Genome: *Homo sapiens*

Tissue: fibroblast

Gene symbol: ITGA2

Type of nucleic acid: mRNA

Invariant bands (kb): 8.0

**Organism:** *Homo sapiens*, human

**Clone type:** Clone

**Shipping information:** Rehydrate with TE

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## Storage Conditions

**Product format:** Frozen

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## Intended Use

This product is intended for laboratory research use only. It is not intended for any animal or human therapeutic use, any human or animal consumption, or any diagnostic use.

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## BSL 1

ATCC determines the biosafety level of a material based on our risk assessment as guided by the current edition of *Biosafety in Microbiological and Biomedical Laboratories (BMBL)*, U.S. Department of Health and Human Services. It is your responsibility to understand the hazards associated with the material per your organization's policies

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and procedures as well as any other applicable regulations as enforced by your local or national agencies.

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**Certificate of Analysis**

For batch-specific test results, refer to the applicable certificate of analysis that can be found at [www.atcc.org](http://www.atcc.org).

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**Insert Information****Insert size (kb):** 5.367**Type of DNA:** cDNA**Insert source:** lung fibroblast IMR-90 cell line**Insert tissue:** lung fibroblast IMR-90 cell line**Insert information:**

DESCRIPTION OF INSERT COMPONENT:

Genomic copy number: unique  
receptor)

Cross references: DNA Seq. Acc.: M28249

Nucleotides 1-5367 of the insert correspond to  
nucleotides 2-5368 of M28249.**Genome:** Homo sapiens**Chromosome:** 5

5 q23-q31

**Gene name:** integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2**Gene product:** integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2 receptor)(collagen  
receptor, platelet gpla/IIa) [ITGA2]**Gene symbol:** ITGA2**Contains complete coding sequence:** Yes**Insert end:** Modification: EcoRI linker**Vector Information**

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**Construct size (kb):** 8.329999923706055**Intact vector size:** 2.964**Vector name:** pBluescript KS+**Type of vector:** phagemid**Construction:** pUC19**Host range:** *Escherichia coli***Vector end:** EcoRI closer to XhoI site; EcoRI**Cloning sites:** SacII; XmaI; NotI; XbaI; SpeI; BamHI; SmaI; PstI; EcoRI; EcoRV; HindIII;

ClaI; SalI; HincII; AclI; XhoI; DraII; ApaI; KpnI

**Enhancer:** none**Insert detection:** lacZ'**Markers:** ampR**Polylinker sites:** SEE COMMENTS**Promoters:** T7; T3; lac**Replicon:** pMB1; f1**Terminator:** none

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## Notes

Restriction digests of the clone give the following sizes (kb): EcoRI--3.6, 3.0, 1.8; HindIII--3.2, 3.2, 2.1; XbaI--8.4; XhoI--4.8, 3.5; KpnI--4.8, 3.5.

- ATCC staff

The insert contains the following restriction sites (approximate kb from the 5' end): EcoRI--1.79, BamHI--1.86; HindIII--2.02, 5.2.

- J. Cell Biol. 109: 397-407, 1989

The coding region is nt 49-3594.

- J. Cell Biol. 109: 397-407, 1989

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## Material Citation

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If use of this material results in a scientific publication, please cite the material in the following manner: clone 2.72F, #19 (ATCC 95491)

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**References**

References and other information relating to this material are available at [www.atcc.org](http://www.atcc.org).

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## Revision

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