



# AQP1 [UTR-01749, WI-9113, pC

## 5675™

### Description

Detects:

Genome: Homo sapiens

Tissue: bone marrow

Gene symbol: AQP1

Type of nucleic acid: mRNA

Invariant bands (kb): 3.1

- **Organism** *Homo sapiens*, human
- **Clone type** Clone
- **Shipping information** Rehydrate with TE

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### Storage Conditions

- **Product format** Dried

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

## 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)** 2.7999999999999998
  - **Type of DNA** cDNA
  - **Insert source** bone marrow
  - **Insert tissue** Bone marrow
  - **Insert information**  
DESCRIPTION OF INSERT COMPONENT:  
Cross references: DNA Seq. Acc.: M77829  
Nucleotides 1-1283 of the insert correspond to nucleotides 53-1335 of M77829.
  - **Genome** Homo sapiens
  - **Chromosome** 7  
7 p14; 7, 84.8799999 cR
  - **Gene name** aquaporin 1 (channel-forming integral protein, 28 kD)
  - **Gene product** aquaporin 1 (channel-forming integral protein, 28 kD) [AQP1]
  - **Gene symbol** AQP1; CHIP28
  - **Contains complete coding sequence** No
  - **Insert end Modification:** EcoRI linker
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## Vector Information

- **Construct size (kb)** 5.760000228881836
- **Intact vector size** 2.961
- **Vector name** pBluescript II KS-
- **Type of vector** phagemid
- **Host range** *Escherichia coli*
- **Vector end** EcoRI
- **Cloning sites** BssHII; KpnI; ApaI; DraII; XhoI; HincII; AccI; Sall; ClaI; HindIII; EcoRI; PstI; SmaI; BamHI; XbaI; NotI; EagI; SacI
- **Insert detection** lacZ', <-, 816-938
- **Markers** ampR
- **MCS** SacI...KpnI, ->, 657-759
- **Polylinker sites** BssHII; KpnI; ApaI; DraII; XhoI; HincII; AccI; Sall; ClaI; HindIII; EcoRI; PstI; SmaI; BamHI; XbaI; NotI; EagI; SacI
- **Primer site** M13-20, ->; reverse, <-

- **Promoters** T3; T7
  - **Replicon** fl, →, 3-459; pMB1, 1032-1972
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## Notes

Restriction digests of the clone give the following sizes (kb): EcoRI--3.0, 2.8; HindIII--4.4, 1.5; SmaI--5.0, 0.8; KpnI--5.8; BglII--5.8.  
- ATCC staff

The insert contains the following restriction sites (approximate kb from the 5' end): SmaI--0.80; HindIII--1.28.  
- GenBank/EMBL/DDBJ

The insert begins at nt +15 relative to the initiation ATG, and contains about 1.3 kb at the 3' end that has not been sequenced.  
- Proc. Natl. Acad. Sci. USA 88: 11110-11114, 1991

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## Material Citation

If use of this material results in a scientific publication, please cite the material in the following manner: pAQP1 [UTR-01749, WI-9113, pCHIP] (ATCC 95675)

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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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