

Datasheet for ABIN1721089

MagSi-S 1.0 beads

2 Images



Overview

Quantity: 2 mL

Application: Separation (Sep)

Product Details

Superparamagnetic silica particles for own development use. Purpose: Bead concentration: 6 - 12 x 10⁹ beads/mL Characteristics: Components: Magnetic silica beads with activated surface Material not included: Buffers and Materials (depending on the application) Magnetic separator for bead separation/collecting Mixer/vortex to homogenize samples and resuspend beads (depending on the application) Bead Ligand: unmodified Bead Matrix: Magnetic Silica particles Bead Size: 1 µm

Application Details

Comment: MagSi-Tools are surface activated magnetic particles, intended for covalent immobilization of proteins (e.g. antibodies, enzymes), peptides, nucleic acids or other molecules of interest.

Different surface modifications and bead sizes allow for choosing the optimal product for the

right molecule to be coupled, and for the intended application. Please take into consideration which groups are available on the ligand for coupling, and try to prevent inactivation or hiding

the active or exposed site of the ligand.

After coupling the molecule of interest (ligand) is coupled to the magnetic particles, the

resulting beads can be used in downstream applications such as:

- Isolating specific target proteins, antibodies, nucleic acids, cells, viruses, etc. (preparative applications)
- Detecting specific target proteins, nucleic acids, cells, viruses, etc. (diagnostic applications)
- Immobilizing enzymes, thereby enhancing stability and minimizing auto-catalysis. Magnetic collection of the particle/enzyme complex allows to remove the enzyme from the reaction, and to reuse it in a new reaction.

Protocol:

Magnetic beads are an ideal tool for immobilizing molecules (proteins, enzymes, antibodies, peptides, nucleic acids, etc.) on a solid phase, to be used for e.g. detecting, enriching, or cleaving specific target molecules. The easy and efficient collection of beads in magnetic fields allows for easy rinsing and removal of excess reagents and ligand after coupling the ligand molecule, as well as easy use in downstream applications. The use of magnetic beads does not require columns or centrifugation steps, and are therefore ideal in high-throughput and automated applications

Bead size

Our magnetic beads come in three sizes, 600 nm, 1 μ m and 3 μ m. 600 nm beads have the advantage of having a larger surface area and the sedimentation time of 600 nm MagSi beads is approximately 4 times slower than that of 1.0 μ m beads. This allows longer incubation times without shaking/mixing, and may be important in automated and other high-throughput applications in which shaking/mixing options are often lacking. MagSi beads with a diameter of 3 μ m have stronger magnetic properties and will separate approximately 4x faster than 600 nm beads under same conditions: approximate separation time is \geq 1 minute using a suitable magnet.

Restrictions:

For Research Use only

Handling

Format:	Liquid	
Concentration:	10 mg/mL	
Buffer:	Stored in 0.05% sodium azide	
Preservative:	Sodium azide	
Precaution of Use:	This product contains sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE which	

should be handled by trained staff only.

Handling Advice:

Store beads in well closed vial and in upright position to prevent drying of the beads since this makes them more difficult to re-suspend.

Do not freeze the product!

Vortex bead suspension well before use.

If you expect iron interference in downstream applications, we strongly advise you to rinse the beads before usage.

Before using the beads it is important to rinse with water or PBS to remove any components that could interfere with your test.

Storage:

4°C

Expiry Date:

12 months

Images

Bead Surface	Chemicals needed	Protein binding	Treatment	Comments
Carboxyl ¹ (COOH)	EDC/NHS	Amine groups (from lysine and/or as unblocked N- termini) Lysine, histidine, cysteine, tyrosine etc.	No treatment needed	Can be used to couple most proteins
Aldehyde (CHO)	Aldehyde/ Amine reaction	Amine groups	No treatment needed	Add reducing agent to stabilize amide bond
Thiol (SH)	Redox reaction ³	Free cysteine	Reduce disulphides under non- denaturing conditions to generate free cysteine.	Useful for proteins containing cysteines. Risk of multiple coupling
Amine ² (NH2)	Gluteral- dehyde	Amine/ aldehyde	No treatment needed	Add reducing agent to stabilize amide bond
Tosyl	None	Sulfhydryl, Amine groups	No treatment needed	Useful for antibodies
Hydrazide	Sodium periodate	Oligosacharide moieties	Oxidize glycoprotein under non- denaturing conditions.	Useful for glycoproteins
Ероку	Adsorption/ reaction support	Lysine, histidine, cysteine, tyrosine etc.	No treatment needed	Useful for enzymes

ion of disulfides with 0.1 M DTE (dithioerythrol); coupling of protein at pH below iso-point; deactivate excess thiel with 20 mM PDEA (2-(2-pyridinyiditho) ethane-amine)/ 1M

Surface activation	Formula	Example Applications	
Silica (stored in 0.05% sodium azide)	Si-OH	- End-users' own application (e.g. functionalization of the MagSi beads)	
Carboxyl (stored in PBS, 0.05% sodium azide)	R-COOH	- Protein and peptide immobilization - Antibody immobilization	
Aldehyde (stored in PBS, 0.05% sodium azide)	R-CHO*	- Protein immobilization	
Amine (stored in 0.05% sodium azide)	R-NH ₂	- Protein immobilization	
Sulfydryl (stored in PBS, 0.05% sodium azide)	R-SH*	- Immobilization via target cysteine groups, coupling to gold surfaces	
Tosyl (stored in DSMO:THF 1:1)	R-S O CH,	- Antibody immobilization - Protein and peptide immobilization	
Hydrazide (stored in PBS, 0.05% sodium azide)	R-CO-N ₂ H ₂	- Glycoprotein immobilization - Protein and peptide immobilization	
Epoxy (stored in DSM0:THF 1:1)	R-CH-CH ₂	- Enzyme immobilization - Protein and peptide immobilization	

^{*} coupling of other organic molecules, such as nucleic acids or carbohydrates, is sible. CHO- and SH-beads have a limited stability, and must be used for coupling ligand within 2-3 weeks after production.

Image 1.

Image 2. Active surfaces and example applications of MagSi-tools