

Identifyn Standards for Optical Microscopy Methods™Identifyn® Primary Antibody**Rabbit anti-Human BRCA1 Recombinant Monoclonal, IgG - (BLR023N)***Cat ID# - PA-RR 000007*Identifyn® Secondary Antibodies**Identifyn® SRM Alexa Fluor™ 647 Affinity Purified Goat Anti-Rabbit, F(ab')₂ (H + L)***Cat ID# - SA 000063***Identifyn® SRM Alexa Fluor™ 555 Affinity Purified Donkey Anti-Mouse, F(ab')₂ (H + L)***Cat ID# - SA 000042*

BRCA1 (Breast Cancer 1) is a protein that plays a crucial role in maintaining the genome's stability and preventing cancer development. BRCA1's primary functions include DNA repair, particularly double-strand breaks that provide function in the homologous recombination pathway.

BRCA1 also significantly regulates the cell cycle, ensuring that cells progress through each phase appropriately. It interacts with various proteins involved in cell cycle checkpoints, helping to prevent cells with damaged DNA from dividing. Additionally, BRCA1 acts as a transcriptional co-regulator, influencing the expression of genes involved in various cellular processes, including DNA repair, cell cycle control, and apoptosis. Finally, BRCA1 is involved in modifying the structure of chromatin (chromatin remodeling); this activity helps regulate gene expression and DNA repair processes.

BRCA1 interacts with numerous proteins, forming complexes involved in the aforementioned cellular processes (DNA repair, cell cycle regulation, transcriptional regulation, and chromatin remodeling). Some fundamental protein-protein interactions of BRCA1 include:

BRCA2: BRCA1 forms a complex with BRCA2, which plays a crucial role in homologous recombination-mediated DNA double-strand break repair.

PALB2 (Partner and Localizer of BRCA2): PALB2 binds to both BRCA1 and BRCA2, facilitating the recruitment of BRCA1-BRCA2 complexes to sites of DNA damage and promoting homologous recombination-mediated repair.

RAD51: BRCA1 interacts with RAD51, an essential protein involved in homologous recombination. This interaction promotes the loading of RAD51 onto single-stranded DNA at sites of DNA damage, facilitating the repair process.

BARD1 (BRCA1-Associated RING Domain protein 1): BARD1 forms a heterodimeric complex with BRCA1, and together they possess E3 ubiquitin ligase activity. This complex plays a role in DNA repair and cell cycle regulation.

CtIP (CtBP-interacting protein): BRCA1 interacts with CtIP, which is involved in the initial processing of DNA double-strand breaks to generate single-stranded DNA ends, a crucial step in homologous recombination-mediated repair.

ATM (Ataxia Telangiectasia Mutated): BRCA1 interacts with ATM, a kinase that plays a central role in the DNA damage response pathway. This interaction is important for activating ATM-mediated signaling in response to DNA damage.

p53: BRCA1 interacts with the tumor suppressor protein p53 and participates in regulating p53-mediated transcriptional responses to DNA damage.

In summary, BRCA1 is essential for maintaining genomic stability and preventing the accumulation of mutations that can lead to cancer development. Mutations in the BRCA1 gene are associated with an increased risk of breast, ovarian, and other cancers, as they impair the protein's ability to carry out its normal functions.

Method

HeLa cells were grown on acid-etched #1.5H cover glass to ~50% confluency. DNA damage was induced via treatment with 2 μ M Etoposide-spiked complete growth media for 24 hours, followed by -20°C methanol fixation and blocked with [Thermo Fisher Fish Serum Blocker cat#37527X3](#). The primary antibody, Rabbit anti-Human BRCA1 Recombinant Monoclonal, IgG, was incubated for 1 hour at room temperature at a dilution of 1:25. Cells were washed 3X in PBS. Identifyn® SRM Alexa Fluor™ 647 Affinity Purified Goat Anti-Rabbit, F(a'')₂ (H + L) was incubated at room temperature for 1 hour at a dilution of 1:1000.

The second primary antibody, [Mitochondria Rabbit Recombinant Antibody \(MTC02,2860R\)](#), was then incubated for 1 hour at room temperature at a dilution of 1:250. Cells were washed 3X in PBS. Identifyn® SRM Alexa Fluor™ 555 Affinity Purified Donkey Anti-Mouse, F(ab')₂ (H + L) was incubated at room temperature for 1 hour at a dilution of 1:1000. Cells were washed 5X in PBS and mounted using [ProLong™ Diamond Antifade Mountant with DAPI cat#P36962](#).

Microscope

Zeiss LSM 980 Confocal **with NIR Detection**, Axio Observer.Z1/7, using a Plan-Apochromat 63X/1.40 oil DIC M27, was used with the following parameters for each dye:

Z Stack - (3D) - Panel A

Z-Stack = 171 Slices (5.1 μ m)

Channels = 3
 Scaling (Per Pixel) = 35.30 nm X 35.30 nm X 30.00 nm
 Image Size (Pixels) = 1566 X 1566
 Image Size (Scaled) = 55.27 μ m X 55.27 μ m
 Bit Depth = 16 Bit
 Image Center Position = X: 73.14 mm, Y: 53.81 mm
 Stage Position = X: 73.14 mm, Y: 53.81 mm
 ROI Center Offset = X: 0.00 μ m, Y: 0.00 μ m

Single Plane (2D) - Z-plane 81 of 171 total - Panel B

Z-Stack = 171 Slices (5.1 μ m) - Panels B & C

Channels = 3
 Scaling (Per Pixel) = 35.30 nm X 35.30 nm X 30.00 nm
 Image Size (Pixels) = 856 X 678
 Image Size (Scaled) = 30.21 μ m X 23.93 μ m
 Bit Depth = 16 Bit

Image Center Position = X: 73.14 mm, Y: 53.81 mm

Stage Position = X: 73.14 mm, Y: 53.81 mm

ROI Center Offset = X: 0.00 μ m, Y: 0.00 μ m

647 -

Reflector = None

Contrast Method = Fluorescence

Pinhole = 5.00 AU / 328 μ m

LSM MBS = MBS 488/561/639, 405/730

LSM SBS = SBS LP 640

Laser Wavelength = 639 nm @1.00%

Laser Blanking = Enabled

Scan Mode = Frame

Scan Zoom = 2.4

Rotation = 0°

Sampling = 2.00

Pixel Time = 0.66 μ s

Frame Time = 7.82 s

LSM Scan Speed = 7

Scan Direction = Bidirectional

Line Step = 1

Averaging = 4

Excitation Wavelength = 653

Emission Wavelength = 668

Effective NA = 1.4

Detection Wavelength = 643-735

Imaging Device = Airyscan

Detector Type = GaAsP-PMT

Detector Gain = 800 V

Detector Offset = 0

Detector Digital Gain = 1.0

LSM Plus Mode = 9.3 (3D, Auto)

555 -

Reflector = None

Contrast Method = Fluorescence

Pinhole = 5.00 AU / 286 μ m

LSM MBS = MBS 488/561/639, 405/730

LSM SBS = SBS LP 550

Laser Wavelength = 561 nm @1.00%

Laser Blanking = Enabled

Scan Mode = Frame

Scan Zoom = 2.4

Rotation = 0°

Sampling = 2.00

Pixel Time = 0.66 μ s

Frame Time = 7.82 s

LSM Scan Speed = 7

Scan Direction = Bidirectional

Line Step = 1

Averaging = 4
Excitation Wavelength = 590
Emission Wavelength = 618
Effective NA = 1.4
Detection Wavelength = 573-627
Imaging Device = Airyscan
Detector Type = GaAsP-PMT
Detector Gain = 800 V
Detector Offset = 0
Detector Digital Gain = 1.0
LSM Plus Mode = 9.4 (3D, Auto)

DAPI -

Reflector = None
Contrast Method = Fluorescence
Pinhole = 5.00 AU / 219 μ m
LMS MBS = MBS 488/561/639, 405/730
LSM SBS = SBS SP 505
Laser Wavelength = 405 nm @0.2%
Laser Blanking = Enabled
Scan Mode = Frame
Scan Zoom = 2.4
Rotation = 0°
Sampling = 2.00
Pixel Time = 0.66 μ s
Frame Time = 7.82 s
LSM Scan Speed = 7
Scan Direction = Bidirectional
Line Step = 1
Averaging = 4
Excitation Wavelength = 353
Emission Wavelength = 465
Effective NA = 1.4
Detection Wavelength = 420-480, 495-499
Imaging Device = Airyscan
Detector Type = GaAsP-PMT
Detector Gain = 800 V
Detector Offset = 0
Detector Digital Gain = 1.0
LSM Plus Mode = 7.9 (3D, Auto)

3D Image Processing - Panel A

3D Volume Projection = Precise
Appearance = Threshold 1% all channels, Ramp 0% all channels, Maximum 100% all channels
Background = Black
Light = Brightness 100%, Azimuth 80°, Elongation -130°, Enabled Directional Lighting, Enabled Tone Mapping
Projection = View Angle 35°, Scale Z 100%, Reverse Z-Axis NOT Selected, Stereo Anaglyph NOT Selected, Camera Separation 5% (deselected), Parallax Shift 0% (deselected)

Clipping Image Process = Clipping planes are introduced to pronounce regions of special interest.

X-Y = Activate was selected, textured opaque was chosen, and a red outline of the clipped perimeter was shown; Clip Back was selected, and Clip Front was NOT selected.

Position of Clip = -13%

Clip X = 0°

Clip Y = 0°

Split View - Panel A

The top-left panel features Rabbit anti-Human BRCA1 Recombinant Monoclonal, IgG, visualized with Identifyn® SRM Alexa Fluor™ 647 Affinity Purified Goat Anti-Rabbit, F(a'')₂ (H + L), the top-right panel features mitochondria visualized with Identifyn® SRM Alexa Fluor™ 555 Affinity Purified Donkey Anti-Mouse, F(ab')₂ (H + L), and the bottom-left panel features DAPI nuclear staining. The bottom-right panel features the merged image of all channels.

3D Image Processing - Panel C

3D Volume Projection = Precise

Appearance = Threshold 4% all channels, Ramp 98% all channels, Maximum 100% all channels

Background = Black

Light = Brightness 100%, Azimuth 80°, Elongation -130°, Enabled Directional Lighting, Enabled Tone Mapping

Projection = View Angle 35°, Scale Z 100%, Reverse Z-Axis NOT Selected, Stereo Anaglyph NOT Selected, Camera Separation 5% (deselected), Parallax Shift 0% (deselected)

Clipping Image Process = Clipping planes are introduced to pronounce regions of special interest.

X-Y = Activate was selected, textured opaque was chosen, and a blue outline of the clipped perimeter was shown; Clip Back was selected, and Clip Front was NOT selected.

Position of Clip = -9%

Clip X = 0°

Clip Y = 0°

Summary

Panel A - This panel presents a three-dimensional volume projection of Rabbit anti-Human BRCA1 Recombinant Monoclonal, IgG. It is counterstained with Identifyn® SRM Alexa Fluor™ 647 secondary antibody (white; the default color, red, has been changed to white in Zeiss Zen software to enhance visual contrast) and Mitochondrial primary antibody labeled with Identifyn® SRM Alexa Fluor™ 555 (orange), along with DAPI (blue), using super-resolution Airyscan imaging. The image features a cut plane on the X-Y axis (red outline), positioned at -13 degrees, with angles X and Y set to 0 degrees (flat) to enhance the visualization of Rabbit anti-Human BRCA1 Recombinant Monoclonal, IgG.

within the nucleus. A region of interest (ROI) is indicated by a white box in Panel A for further investigation in Panels B and C.

Panel B - This panel displays a split view from the ROI in Panel A, at a z-plane 81 out of 171, illustrating the signal and localization of Rabbit anti-HumanBRCA1 Recombinant Monoclonal, IgG. The upper left section highlights the primary antibody, Rabbit anti-Human BRCA1 Recombinant Monoclonal, IgG, labeled with Identifyn® SRM Alexa Fluor™ 647 (white; the default color, red, has been changed to white in Zeiss Zen software to enhance visual contrast) The upper right panel shows a Mitochondrial primary antibody labeled with Identifyn® SRM Alexa Fluor™ 555 (orange. The lower left section presents DAPI nuclear staining (blue), while the lower right section features a merged image of the three channels.

Panel C - This panel provides a three-dimensional volume projection of Rabbit anti-Human BRCA1 Recombinant Monoclonal, IgG. It is counterstained with Identifyn® SRM Alexa Fluor™ 647 secondary antibody (white; the default color, red, has been changed to white in Zeiss Zen software to enhance visual contrast), as well as DAPI (blue) at super resolution Airyscan.

Note - At Airyscan resolution, the previously observed "pan-nuclear" or diffuse nature of BRCA1, seen in diffraction-limited techniques such as widefield imaging, widefield structured illumination microscopy (SIM), and confocal microscopy, now reveals enhanced structural details. Research on nuclear proteins using Identifyn's™ step-down method indicates that what was once thought to be ubiquitous or pan-nuclear now appears interconnected and organized. Further, Airyscan technology illustrates that BRCA1 is localized outside of the nucleolus, as expected, and additionally reveals both a strong localized signal and some "thread-like" structure. While further research is needed to determine the exact localization and meaning of these structures, it is evident that as resolution decreases, more information becomes accessible to researchers.

Image Corrections

Identifyn® SRM Alexa Fluor™ 647 Affinity Purified Goat Anti-Rabbit, F(ab')₂ (H + L) was pseudo-colored in Zen Software from red, the default color, to white to provide visual contrast.

Image by J.K. Bennett
Image Analysis by B.T. Bennett
Data Presentation by P.D. Amistoso

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