Biodynamic optical assay for embryo viability

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Abstract. Early stage porcine parthenogenetic embryos were evaluated for metabolic activity using a biodynamic microscope (BDM) that images dynamic light scattering using low-coherence digital holography. The microscope has a 45-deg illumination configuration that reduces specular background for the imaging of small translucent samples. The off-axis illumination is compatible with coherence-gated imaging because of volumetric light scattering in which the coherence plane is tilted at half the illumination angle in a three-dimensional tissue target. The BDM was used to profile the viability of porcine parthenotes with normal and with inhibited mitochondrial adenosine triphosphate (ATP) production using Doppler fluctuation spectroscopy. The ATP concentrations in the parthenotes, which are indicative of developmental potential, were validated by a conventional bioluminescence assay. Biodynamic classifications achieved ~80% accuracy correlating sample ATP treatment, providing a quick, label-free surrogate measurement to replace invasive metabolic assays as a candidate for evaluating quality of early embryos in the assisted reproductive technology setting.

In the United States, 7% of married women aged 15–44 are infertile, and in-vitro fertilization (IVF) technology is growing at about 5% per year in terms of IVF cycles. The IVF birth success over all ages of women was only about 30% in 2016.1,2 The IVF success rates are connected to the viability of the blastocyst prior to transfer in utero, and blastocyst quality is related to implantation selection and pregnancy potential.3 Because of the low success rates, clinics typically transfer two or three embryos at the same time. However, this can lead to multiple pregnancy and increases the risk of adverse health conditions for both the mother and the offspring. The best practice is to transfer a single high-quality embryo. Morphological grading, using different grading systems, is used to evaluate embryo quality, but it is a subjective process.4 Other noninvasive methods include analysis of spent culture media, metabolomics, time-lapse monitoring of embryo development,5 and dynamic index variations from quantitative phase imaging methods have shown potential to predict embryo viability.6 However, there is no reliable evidence from randomized controlled trials to show that these noninvasive techniques are effective in improving the birth rate.7,8 Invasive methods may involve biopsy or manipulation of blastocysts9 and may be detrimental to later development of samples.10 In this letter, we describe the application of biodynamic imaging (BDI) to the problem of viable embryo selection. Biodynamic imaging based on low-coherence Doppler fluctuation spectroscopy,11 which captures dynamic activity dominated by intracellular transport, has been used successfully to profile cumulus-oocyte complexes (COCs).12 Drug response of three-dimensional (3-D) cultured tumor spheroids,13,14 and to classify cancer patients in preclinical trials.15 In this letter, BDI is demonstrated as an accurate surrogate for metabolic assays in embryo selection.

The samples used to perform this study were porcine parthenotes. Parthenotes are embryos derived from oocytes that are activated parthenogenetically by stimuli such as electric shock or exposure to chemicals, which causes them to resume meiosis and stimulates their development. Pig parthenotes undergo cell divisions and can form blastocysts (but are unable to develop to term). Normally, embryos originate from both the male and female gametes that fuse at fertilization, but a parthenote develops from an oocyte only. Nevertheless, parthenotes are frequently used in developmental studies in the laboratory because they behave similarly to IVF embryos.16

The biodynamic microscope (BDM) is a stand-alone module, manufactured by Animated Dynamics, Inc., that inserts into a conventional Olympus IX-73 inverted microscope, allowing a user to switch between conventional transillumination microscopy and a BDI mode. The BDM interferometry system is a Mach–Zehnder configuration. A Superlum™ superluminescent diode centered at 841.2 nm with a 28.9-nm bandwidth and 20.9-mW output power was used as the light source that was incident on the sample at a 45-deg angle relative to the optic axis of the collection objective lens. The translation stage controls the coherence gate based on optical path length (OPL). In conventional backscatter coherence-gated holography, the optical section plane is perpendicular to the optic axis. However, the 45-deg illumination, combined with the three-dimensional volumetric scattering, creates a coherence plane that is tilted at 22.5 deg (half of the illumination angle), and the sample “flythrough” occurs at this oblique angle when translating the reference mirrors. The advantages of this BDM design are the reduction of specular background reflections and the elimination of the conventional beamsplitter normally used in the backscatter configuration to improve optical brightness of dim translucent samples. The oblique coherence plane needs minor additional digital postprocessing after holographic reconstruction to regain a balanced aspect ratio of the volumetric target.

The optical configuration is shown in Figs. 1(a) and 1(b). The optical design of the BDM uses an Olympus UPLFLN 4× objective lens with a 0.13-numerical aperture, 17-mm working distance, and 45-mm parfocal distance. The BDI mode has a 0.05-numerical aperture, up to 1-mm field of view (determined by the illumination beam size), and a pixel size of 8 μm. The BDM is placed on a vibration isolation platform to minimize the mechanical disturbance.
Ovaries were collected from slaughtered pigs, and follicular contents were aspirated from individual follicles in the lab using a hypodermic needle attached to a 10-mL syringe. The oocytes with attached cumulus cells (COCs) were collected from the follicular fluid by means of a stereomicroscope, washed in TL-HEPES medium, and good-quality oocytes were selected for in-vitro maturation in the appropriate medium. Matured COCs were placed in 0.1% hyaluronidase, vortexed until denuded and washed in TL-HEPES medium. Denuded oocytes were then placed in electroreporation medium in a chamber containing two stainless steel electrodes 0.5-mm apart. Parthenogenetic development was induced via electroreporation with two direct-current pulses 1-s apart of 1.2 kV/cm, 60 μs each via a CF-150/B cell fusion instrument. After electric stimulation, prospective embryos were rinsed in TL-HEPES medium and placed in 50-μL droplets of PZM-3 medium with or without 20-μM sodium azide (NaN₃). When an embryo is cultured with dilute sodium azide (NaN₃) at the 1-cell stage, the azide ion (N₃⁻) inhibits the electron transport in the mitochondrial membrane and prevents oxidative phosphorylation. Mitochondrial adenosine triphosphate (ATP) production of the embryo is thus inhibited, and its ability to develop into a healthy blastocyst is significantly reduced.¹⁷,¹⁸ Each droplet contained 10 embryos, the droplets were covered with light mineral oil to prevent evaporation, and culture dishes containing the droplets were placed in a CO₂ incubator. Both groups were allowed to culture in their respective medium up to 96 h, at 39°C, under 5% CO₂ in air.

After the culture period, the embryos that had reached the early morula stage were selected for assessment. For BDI measurements, samples were prepared with two protocols with different immobilization methods: (1) each sample was prepared in a 50-μL TL-HEPES droplet on a plate treated with Cell-Tak™ and covered with mineral oil, or (2) three washing dishes of TL-HEPES medium, and good-quality oocytes were selected for maturation in the appropriate medium. Matured COCs were rinsed in TL-HEPES medium and good-quality oocytes were selected for in-vitro maturation in the appropriate medium. Matured COCs were placed in 0.1% hyaluronidase, vortexed until denuded and washed in TL-HEPES medium. Denuded oocytes were then placed in electroreporation medium in a chamber containing two stainless steel electrodes 0.5-mm apart. Parthenogenetic development was induced via electroreporation with two direct-current pulses 1-s apart of 1.2 kV/cm, 60 μs each via a CF-150/B cell fusion instrument. After electric stimulation, prospective embryos were rinsed in TL-HEPES medium and placed in 50-μL droplets of PZM-3 medium with or without 20-μM sodium azide (NaN₃). When an embryo is cultured with dilute sodium azide (NaN₃) at the 1-cell stage, the azide ion (N₃⁻) inhibits the electron transport in the mitochondrial membrane and prevents oxidative phosphorylation. Mitochondrial adenosine triphosphate (ATP) production of the embryo is thus inhibited, and its ability to develop into a healthy blastocyst is significantly reduced.¹⁷,¹⁸ Each droplet contained 10 embryos, the droplets were covered with light mineral oil to prevent evaporation, and culture dishes containing the droplets were placed in a CO₂ incubator. Both groups were allowed to culture in their respective medium up to 96 h, at 39°C, under 5% CO₂ in air.

Samples are assessed using the BDM by evaluating speckle fluctuation properties. The average backscatter brightness (BB, i.e., intensity) and normalized standard deviation (NSD) for each pixel are calculated over 500 frames (NSD is defined as ΔI/I, where ΔI is the standard deviation of intensity and I is average intensity). Each sample is characterized by optical coherence imaging (OCI) shown in Fig. 2(b), by motility contrast imaging (MCI) shown in Fig. 2(c), and by the Doppler power spectrum (DPS) shown in Figs. 2(d) and 2(e). Sample spectra were prepared, and each sample was washed in each of the dishes and plated in 50-μL PVA-free TL-HEPES droplet and covered with mineral oil. The plate was heated with the temperature maintained at a physiological 39°C. A complete BDI dataset for a sample contained 100 background frames and 2500 holograms, captured at 25 fps with an exposure time of 20 ms, reconstructed with Fourier transform [Fig. 1(c)]. The coherence gate was placed approximately at the middle of the sample at a depth of about 100 μm from the embryo surface. The BDI measurement finished in a short time (1 to 2 min of sample preparation and 2 min of data acquisition), by placing the samples within a safe light exposure range and a non-CO₂-controlled environment. After the BDI measurement, each embryo was lysed in 10 μL of RIPA lysis and extraction buffer and stored for 10 min on ice. ATP measurements were then performed using Molecular Probes™ ATP Determination Kit on a Spark™ 10M multimode microplate reader. A standard curve was obtained based on bioluminescence from the reactions of standard solutions with known ATP amount. Bioluminescence of each sample was measured and compared against the standard curve and converted to the respective ATP amount of substance. A total of 133 embryo samples measured on 12 separate days were used in the analysis, consisting of 85 control samples and 48 NaN₃-treated samples. The BDI data for the samples with different immobilization methods were normalized and combined in the analysis.
span a range of 0.01 to 2 Hz and are fit with a “stretched” Lorentzian lineshape
\[
S = \frac{A}{\omega_0^2 + \omega^2} + N_y,
\]
where \( \omega_0 \) is the knee frequency (the “roll-over” frequency of the spectrum, i.e., where the curvature of the curve changes), \( s \) is the slope in the midfrequency range, and \( N_y \) is the Nyquist floor. These parameters are the potential biomarkers of embryo viability. Typically, an azide-treated spectrum displays a lower knee frequency, which is correlated with slower intracellular activities. Azide-treated spectra also have a greater dynamic range (DR) and a steeper midfrequency slope on a log-log power spectrum graph. In addition, the spectrum is characterized by the slope and \( R^2 \) values from linear fittings on the log scale (i.e., a power law fitting model \( S = a \omega^{-b} \)), both “globally” on the entire spectrum and “locally” in three frequency ranges: 0.01 to 0.08 Hz, 0.08 to 0.4 Hz, and 0.4 to 2 Hz. Values of \( b \) and \( R^2 \) are used to describe the spectrum shape (values for the three ranges use the notations \( b_1, b_2, b_1, r_1, r_2, \) and \( r_3 \)). Typical values for these parameters across a range of parthenotes are listed in Table 1.

![Fig. 2](image)

**Fig. 2** (a) Microscope image of a parthenote sample captured in transillumination. (b) OCI of the sample. Color map is on log scale. (c) MCI of the same sample. The value of each pixel is NSD (scale bar = 100 \( \mu \)m in a, b, and c). (d) Averaged fluctuation spectra (smoothed) of control samples and NaN3-treated samples plated in PVA-free TL-HEPES droplets, with different knee frequencies, slopes, Nyquist floors, and dynamic ranges (“DR”) on the log-log scale. Standard errors are used as values of error bars. (e) An example of three-segment linear fitting for spectra. (Markers are a subset of data points with even intervals.)

Analysis of the sample ATP content shows that the control group parthenotes have significantly more ATP than for these parameters across a range of parthenotes are listed in Table 1. ATP content greater than a threshold as the varying parameter as shown in Fig. 3(b). This classifier achieved 89.7% accuracy at an ATP threshold of 5 nmol.

In this correlative study, biomarkers were used to build sample classifications and to estimate the predictive ability of the BDM assay to select embryos with high metabolic activity. A feature vector contains the key biodynamic biomarkers that represent sample properties with 13 elements \( \{ \text{BB, NSD, knee, midfrequency slope, } R^2 \text{ value of spectrum, floor, DR, and } b \text{ and } R^2 \text{ values from three frequency ranges}\}. \) Principal component analysis (PCA) was used for dimensionality reduction, after which a quadratic-kernel support vector machine (SVM) was used to predict the sample group (using MATLAB® Statistics and Machine Learning Toolbox™). In this analysis, the “control” group was defined as condition positive, while the NaN3-treated samples were defined as condition negative. A fivefold cross validation was used to characterize the performance of the classifier, where a classification score, indicating a signed distance from the observation to the decision boundary, was assigned to each sample. Choosing a fivefold cross validation ensured that the test groups were large enough (with \( n \sim 26 \) in each group) and that the bias was low. The fivefold cross validation was run 10 times, and in each run TPR and FPR values were calculated with varying score thresholds, creating the receiver operating characteristic (ROC) curves plotted in Fig. 3(c). The SVM classifier performed with an area under the curve (AUC) of 0.812 and an accuracy of 79.3%.

In conclusion, BDI of intracellular activity has been demonstrated on early stage parthenotes as biologically relevant models of natural embryos. The biodynamic assay performs as a surrogate for invasive ATP assays and can distinguish parthenotes that have high metabolic activity from parthenotes that have compromised metabolism. This assay is noninvasive and can be performed longitudinally to track embryo health while preserving embryo viability. This optical technique has the

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<th>Table 1 Typical values for biodynamic biomarkers.</th>
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<td><strong>Quantity</strong></td>
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potential to improve IVF success rates. Future studies will include a prospective preclinical trial during which porcine embryos will be evaluated with BDI and then transferred in utero in sows to establish pregnancy rates for BDI-selected transfer relative to conventional selection.

Disclosures
Professor Nolte has a financial interest in Animated Dynamics, Inc., that provided the biodynamic microscope used for this project, and Professor Machaty is a paid consultant.

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References

Fig. 3 (a) Box plot of ATP content in “control” parthenotes and NaN3-treated parthenotes. (b) ROC curve of the sample ATP biochemical assay and (c) ROC curves and averages from 10 runs of the fivefold cross-validated SVM classifier on the biodynamic feature vector. The blue curve is the average of TPRs for each FPR, while the gray area is the ±1 standard deviation of the TPRs.