# Cytokines and chemokines in follicular fluids and potential of the corresponding embryo: the role of granulocyte colony-stimulating factor

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**BACKGROUND:** The cytokine/chemokine levels of individual follicular fluids (FFs) were measured to determine whether a biomarker could be linked to the developmental potential of the derived embryo. **METHODS:** Fluid was collected from 132 individual FFs that were the source of oocytes subsequently fertilized and transferred. In each, a bead-based multiplex sandwich immunoassay (Luminex) was used to measure 28 cytokines and chemokines simultaneously. **RESULTS:** Significantly higher levels of interleukin (IL-2) and interferon (IFN-γ) were detected in FF for embryos that underwent early cleavage. IL-12 was significantly higher in FF corresponding to highly fragmented embryos and the chemokine CCL5 was significantly higher in FF related to the best quality (Top) embryos. The level of granulocyte colony-stimulating factor (G-CSF) in individual FF samples was correlated with the implantation potential of the corresponding embryo. The area under the receiver operating characteristics curve, which distinguished the embryos that definitely led to delivery from those that did not, was 0.84 (0.75-0.90) (P = 0.0001) for FF G-CSF. FF G-CSF was significantly lower in patients older than 36 years compared with those <30-year old. When the FF G-CSF was 20 pg/ml or higher, the ratio between Top and non-Top embryos was significantly higher than for the group with FF G-CSF below 20 pg/ml (45 versus 20.45%, P = 0.007). **CONCLUSIONS:** Individual FF composition is related to the development of the corresponding *in vitro* generated embryo and its potential of implantation. Individual FF G-CSF may provide a non-invasive biomarker of implantation that needs to be evaluated together with *in vitro* observation to select the oocyte, and hence the embryo, to transfer.

Keywords: G-CSF; follicular fluid; embryo implantation; IVF/ICSI-ET; bead-based multiplex immunoassays

# Introduction

Assessing the implantation potential of an embryo to be transferred is crucial for increasing the success rates of IVF-ICSI cycles while reducing or eliminating the risk of multiple pregnancies. The application of the 'single embryo transfer' (SET) policy has proven effective in decreasing the maternal and foetal morbidity and mortality associated with assisted reproductive technologies (De Neubourg and Gerris, 2003; De Sutter *et al.*, 2003; Gerris *et al.*, 2004; Pinborg, 2005; Fiddelers *et al.*, 2006). This policy, however, is applied only sporadically worldwide, except when imposed by a specific legislation in a given country, mainly because the mean embryo implantation rates remain too low (15-20%). The transfer of at least two embryos significantly increases overall pregnancy rates (van Montfoort *et al.*, 2006), and two embryos are routinely transferred into the uterine cavity in Europe, and 3-4 in the USA. The application of SET would be encouraged and the risk of multiple pregnancies minimized in IVF-ICSI cycles if methods could be found to select those embryos with the highest implantation potential so as to transfer them selectively.

The analysis of the morphology of the pre-implantation embryo before Day 2, although important, is generally not sufficiently informative. Additional data about the embryo's potential are required (Guerif *et al.*, 2007). An alternative approach to applying SET would be to select the best oocyte and thereby avoid the ethical problems linked to the generation of surplus embryos and comply with the law in countries where all fertilized embryos must be transferred. Identifying prognostic markers of oocyte competence is difficult, but cytokines and chemokines present in individual follicular fluid (FF), which surrounds the oocyte-cumulus oophorus complex, may prove useful. Several previous studies have described factors in FF that appear to be related to a successful outcome in IVF-ET (i.e. a live birth) (Hammadeh *et al.*, 2002; Ocal *et al.*, 2004; Westergaard *et al.*, 2004; Wiener-Megnazi *et al.*, 2004; Asimakopoulos *et al.*, 2005; Salmassi *et al.*, 2005; Rizzo *et al.*, 2007). However, these studies have generally been performed with pooled FF in which fluids from functional cysts, atretic, immature and mature follicles were all mixed. Studies of FF that is pooled or of limited traceability (Fanchin *et al.*, 2007) cannot demonstrate whether detection of these factors allows the selection of high-quality oocytes on an individual basis. To the best of our knowledge, however, no marker in an individual FF sample is known to be

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related to the implantation potential of the resulting embryo after a conventional ovarian stimulation protocol.

We therefore decided to use multiplex analysis to explore the patterns of expression of cytokines, chemokines and growth factors of the FF of individual oocytes and then to relate these patterns to the morphology and implantation potential of the embryo derived from the oocyte from that follicle.

To maximize the potential of the multiplex assay, we limited our targets to molecules either known to participate in the early embryonic 'immunotrophic' activity or suspected of involvement in preparing local (uterine) tolerance to the conceptus. We therefore focused on those cytokines which are most likely to be involved in switching the local Th-1/Th-2 equilibrium towards a Th2 profile (Wegmann *et al.*, 1993). We also studied the TNF family, its related chemokines and the growth factors which are generally believed to be involved in mobilizing the surrounding immune cells. The main outcome measure of the study was the live delivery of each corresponding embryo.

We report here that G-CSF in individual FF appears to be an immune biomarker of implantation, related to ageing and ovarian stimulation and providing information complementary to that of embryo morphology.

## Study protocol

### Patients

We recruited 142 women being treated for infertility and included in the ICSI program at the Obstetrics and Gynaecology department of Poissy-St Germain en Laye Hospital. To avoid any possibility of a bias in patient selection, the patients were randomly allocated (1/1) at inclusion in the ICSI program into either a group where individual FF were collected at oocyte retrieval (study group n = 71) or a group with standard oocyte retrieval where FF were pooled (n = 71). Each patient was included only once during the study period.

All patients were fully informed, and the Institutional Review Board (Comité Consultatif de Protection des Personnes, Poissy-St Germain en Laye) approved this investigation.

The mean age of the population was 33 years (range: 24-41), the mean number of previous IVF attempts was 2 (1-6) and the mean time since first attempting conception was 4 years (1-12 years). In 49 cases, the infertility was primary and in 22 cases, it was secondary. In 56 cases, the infertility was completely attributable to the man, in 10, it was mixed and in 5, it was unexplained, with a previous IVF failure.

# Treatment

The ovarian stimulation protocols were selected by each treating physician: a standard long protocol with a daily GnRH agonist for 54 patients, a short protocol with a daily GnRH agonist for 11 patients and an antagonist protocol for six patients. The response to stimulation was monitored by serial blood tests as well as by ultrasound assessment of follicular and endometrial growth. Ovulation was triggered when at least four follicles had reached 16 mm. We classified the responses to ovarian stimulation in three categories: low response (estradiol (E2) below 1500 pg/ml the day ovulation was triggered), mild response (E2 from 1500 through 3500 pg/ml) and high response (E2 over 3500 pg/ml).

Oocytes were retrieved by aspiration 35-36 h after the triggering of ovulation, with a general or local anaesthesia and vaginal ultrasound guidance. An individual 10-ml syringe was used for each follicle in the study group. We thus adapted the standard method of oocyte aspiration to collect the FF of each individual oocyte separately.

# FF samples

Individual FF samples, each corresponding to one mature oocyte, were collected from women in the study group. The presence of an extra-technician was necessary at the time of the collection to record the presence of an oocyte in the FF sample collected. Samples without an oocyte were immediately discarded. The volume and appearance (yellow or reddish and bloody) of each FF sample were recorded. Each sample was centrifuged and divided into aliquots after anonymization of the samples in the database. This ensured that the subsequent analyses were performed on a blinded basis.

The samples were initially stored at -20°C and then at -80°C until they were assayed.

Only the FF samples corresponding to embryos actually transferred on Day 2 were analysed. Accordingly, of 530 FF samples collected, 146 embryos were transferred, and the fluid from follicles related to 132 of them was analysed. For 15 patients in a sub-cohort, FF retrieved from every follicle aspirated during the cycle was analysed (76 samples) to study the individual variations.

Oocyte fertilization and embryo culture until Day 2

Oocytes were collected and the cumulus and corona cells removed with hyaluronidase 80 IU (Fertipro, JCD, France). The oocytes were injected with single sperm in a 5-µl droplet of Ferticult Hepes (JCD, France), with

viscous PVP medium (Fertipro, JCD, France) to slow the sperm. The injected oocytes were individually cultured in a 40-µl micro-droplet of ISM1 (Medicult, France) under oil at 37°C in a 5% CO<sub>2</sub> humid atmosphere. The number of pronuclei and their appearance were assessed by Gianaroli's criteria after 20 h (Gianaroli *et al.*, 2003). At 25 h, we assessed the early cleavage rate. On Day 2, we assessed and recorded the number, fragmentation and regularity of each individual blastomere. The decision to transfer an embryo was taken by the embryologist in charge, taking in account the morphology at 18 h, 25 h and Day 2 and the rate of development during the first 2 days. Embryo transfer was scheduled on Day 2. The transferred embryos were observed. (i) At the zygote stage (18-20 h), the pronuclear and nuclear morphology and polar body alignment were recorded for 76 embryos according to the classification described previously (Gianaroli, 2003). (ii) At 25 h, early cleavage was recorded for 92 embryos. (iii) On Day 2, the embryos were analysed in relation to:

- (a) Their fragmentation (Grade 1: <10%, Grade 2: 10-30%, Grade 3: 30-50%, Grade 4: over 50% (the corresponding embryo was downgraded when blastomeres were unequal in size)
- (b) Their embryo score: number of blastomeres  $\times$  (5 embryo grade) (Steer *et al.*, 1992). We created three categories of analysis: score 16-30, 15-12 and <12.
- (c) According to the 'Top' versus 'not Top' analysis: 'Top' embryos were defined as those with 4-5 cells on Day 2, <10% fragmentation and regular cells. All other embryos were classified as 'not Top'.

Determination of cytokine and chemokine concentrations in individual FF samples with bead-based multiplex sandwich immunoassays and ELISA measurements

We used a bead-based multiplex sandwich immunoassay, read with a Luminex system (Luminex Map Technology) to measure the concentrations in individual FF samples of the following cytokines and chemokines: IL-1 $\alpha$ , IL-1Ra, IL-2, IL-4, IL-5, IL-6, IL-8, IL-9, IL-10, IL-12, IL-13, IL-15, IL-17, IFN- $\alpha$ , TNF- $\alpha$ , G-CSF, GM-CSF, VEGF, PDGF, FGF, IP-10, MCP-1, CCL5, eotaxin, MIP-1- $\alpha$  and MIP-1- $\beta$ . In brief, 50  $\mu$ l of each individual FF sample was added to 50  $\mu$ l of antibody-conjugated beads directed against the cytokines listed above (Bio-Rad Laboratories, Hercules, CA, USA) in a 96-well filter plate (Bio-Rad). After a 30-min incubation, the plate was washed and 25  $\mu$ l of biotinylated anti-cytokine antibody solution was added to each well before another 30-min incubation. The plate was then washed and 50  $\mu$ l of streptavidin-conjugated PE was added to each well. After a final wash, each well was resuspended with 125  $\mu$ l of assay buffer (Bio-Rad) and analysed by the Luminex array system. The cytokine concentrations were calculated and a standard curve was derived from various concentrations of a cytokine standard in the assay.

The limits of detection for G-CSF for the Bio-Rad kit are  $5.33 \pm 9.12$  pg/ml (mean  $\pm 3 \times$  SD of the blanks, where G-CSF was not present). The intratest variation was assessed by measuring the same FF samples 10 times and the intratest SD for G-CSF ranged from 0.18 to 0.79 pg/ml.

As reagents for measuring LIF are not available for the Luminex system, it was analysed with the non-commercial ELISA previously described (Taupin *et al.*, 1997), which combines two monoclonal antibodies known to not interfere with ligand receptors.

Evaluation of implantation and delivery potential

The probability of implantation, described here as the clinical implantation rate (IR), was defined for each sample tested as the ratio of the number of intrauterine gestational sacs to the number of transferred embryos. Clinical implantation was defined at 8 weeks of amenorrhea when a gestational sac was viewed by ultrasound. The delivery rate was defined as the ratio of the number of babies born to the number of embryos transferred.

The 'certain implantation and delivery' category for receiver operating characteristics (ROC) curve analysis was defined when the number of embryos replaced was equal to the number of gestational sacs or babies born (i.e. one embryo replaced and a singleton pregnancy, two embryos replaced and a twin pregnancy).

Statistical analysis

In order to explore the significance of the concentrations of cytokines and growth factors in each FF with regard to the clinical profile and response of the stimulation of each patient, we created categories of analysis and used the ANOVA test to highlight significant differences.

The following demographic data were analysed:

- (i) Age (younger than 30 years old, from 30 to 36, 37 years or older)
- (ii) Type of infertility (primary or secondary)
- (iii) Aetiology
- (iv) Number of previous attempts (1 or 2, 3 or 4, over 4)
- (v) Ovarian reserve, based on the hormone assays performed on Day 3 of the cycle.

The basal FSH was defined as elevated over 9 IU/ml and normal below 9 IU/ml.

- (i) Protocol type (long or short protocols or antagonist)
- (ii) Hormonal response to stimulation measured by the  $E_2$  level on the day of the HCG injection (1500 pg/ml or less, from 1500 through 3000, more than 3000).

For the embryo morphology analysis, we took into account at 20 h the pronuclear and nuclear morphology and the polar body alignment at 20 h, the occurrence of early cleavage at 25 h, grading according to the fragmentation, embryo score, and classification (Top versus not Top at 48 h).

To construct the ROC curves, we took into account only two categories: 'no delivery' and 'definite delivery' (when the number of embryos replaced equalled the number of babies born). The discrimination between no and definite delivery as a function of the concentration of each cytokine and chemokine in each sample was evaluated with ROC curve analysis (MedCalc for Windows, version 9.2.0.0 software, MedCalc Software, Mariakerke, Belgium). In a ROC curve the true positive rate (sensitivity) is plotted as a function of the false positive rate (100-specificity) for different cut-off points (Zweig and Campbell, 1993). The calculation of the area under the ROC (AUC-ROC) curve measures the accuracy, i.e. the ability of the cytokines and chemokines tested to discriminate between a successful and a failed implantation. The ROC curve analysis thus allowed us to define the lower limit implantation probability, defined by a negative predictive value of 100% from the AUC-ROC curve, and its upper limit, defined by the highest positive predictive value for implantation.

We were therefore able to compare the implantation and delivery potentials of each embryo as a function of the G-CSF concentration in a predefined category. A value of P < 0.05 was considered significant.

### Results

ICSI results and pregnancy/delivery rates in the observed cohort

The mean number of oocytes collected was 8, the mean number of embryos obtained 4.5, and the mean number of embryos transferred 2 (1-3). The mean implantation rate per embryo was 21% (31 gestational sacs for 146 transferred embryos), the mean clinical pregnancy rate 30.9% (22/71) and the mean live birth rate 29.6% (21/71). The multiple pregnancy rate was 36% (8/22).

Detection of cytokines and chemokines

LIF, IL-1Ra, IL-4, IL-6, IL-8, IL-10, IL-12, IL-13, G-CSF, VEGF, IP-10, MCP-1, eotaxin and MIP- $\beta$  were detected in all FF samples, while IL-1- $\beta$ , IL-5, IL-7, IL-17, TNF- $\alpha$  and MIP- $\alpha$  were detected in none. IL-15, GM-CSF, CCL5, PDGF, IFN-7, IL-9, IL-2 and FGF were detected, respectively, in 95, 88, 81, 76, 65, 60, 48 and 22% of the FF samples (Table I).

Variations of the cytokine and chemokine content of FF in relation to demographic data

FF samples of 38 follicles were collected from patients younger than 30 years old, 57 from patients between 31 and 36, 57 over 37 years old. We observed a decrease in FF G-CSF with ageing, which was, respectively, 22.2, 21.3 and 19.6 pg/ml (Fig. 1). FF from patient younger than 30 years had a significant higher G-CSF content than those from patients 37-year old or more (P = 0.03).

We did not observe any significant differences according to the range of the ICSI attempt for the cytokines and growth factors tested.

FF G-CSF and IL-13 levels were significantly higher in patients with primary compared to those with secondary infertility (21.6 and 4.6 versus 19.9 and 4.2 pg/ml, respectively, P = 0.04 and 0.003), while CCL5 levels were significantly lower (321.56 versus 733 pg/ml, P = 0.01).

FF IL-15 was significantly higher in patients with unexplained infertility compared to those with male or mixed infertility (5 versus 1.3 and 1.6 pg/ml for IL-15, P = 0.01 and 0.03, respectively).

Hormonal response to ovarian stimulation and cytokine/chemokine concentrations in individual FF samples

FF samples were classified according to hormonal response to stimulation: low (n = 23), normal (n = 90) and high responders (n = 9) were identified. The level of G-CSF was significantly lower in high compared with low responders (18.5 versus 22.4 pg/ml, P = 0.02). We observed significant differences for many FF cytokines (G-CSF, IL-8, IL-15, IL-17, IFN-7, VEGF, IP-10, CCL5 and MIP- $\beta$ ), but always in the same direction, that is, the FF cytokine levels decreased when  $E_2$  levels increased.

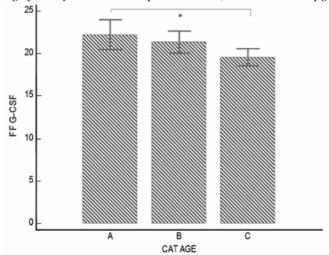
No significant difference was observed according to the stimulation protocol used.

**Table I.** Mean values, standard error and standard variation of the cytokines and chemokines detected in individual FF samples with the Luminex technology (bead-based multiplex sandwich immunoassays).

Cytokines/chemokines (pg/ml)	Mean	Standard deviation	Standard error
IL-1Ra	225	530	46
IL-2	8	5.8	0.5
IL-4	1.8	0.7	0.06
IL-6	21.2	79	6.8
IL-8	399	2785	241
IL-9	9.9	13.4	1.16
IL-10	4.6	4.6	0.4
IL-12	15.3	6.2	0.53
IL-13	4.5	0.73	0.064
IL-15	1.77	3.76	0.32
IFN-γ	32.9	43.1	3.7
G-CSF	21.06	4.64	0.40
GM-CSF	25.4	11.1	0.96
VEGF	12 616	13 565	1176
PDGF	248.8	1388	120
FGF	19	47.6	4.1
IP-10	2083	1948	168.9
MCP-1	256.6	1560	135
CCL5	449	1087	94
Eotaxin	138	103	8.9
MIP-1 beta	266	1989	172
LIF	954	1150	103

Those shown in bold have acceptable standard deviations and errors for the means.

**Figure 1:** Effect on ageing on individual FF G-CSF concentration (pg/ml). Three groups are shown: (A) <30-year old (n = 38), (B) between 30 and 36-year old (n = 57), (C) older than 36 years (n = 57). \*FF G-CSF is significantly lower in Group A versus C (22.2 versus 19.6 pg/ml, P = 0.03).



Cytokine and chemokine concentrations and the appearance of FF

FFs were classified according to the appearance and volume of each sample at the time of the collection in order to evaluate the impact of blood contamination on the cytokines/chemokines content or follicle volume. In all, 33 were yellow, 58 reddish and 48 bloody, 31 had a volume of 2 ml, 92 had a volume between 3 and 6 ml and 10 had a volume over 7 ml. The cytokine/chemokine content did not vary according to the appearance or volume of the fluid collected.

Cytokine and chemokine concentrations in individual FFs and subsequent morphology of the transferred embryos

Among the 78 embryos observed at the zygote stage, cytokine or chemokine content did not differ according to categories of pronuclear and nuclear embryo staging. There were 33 embryos with a longitudinal polar body

alignment (alpha group), 28 with a perpendicular alignment (beta group) and 19 other patterns (gamma group). IL-1Ra and TNF were significantly elevated in the gamma group compared with the alpha and beta groups (485 versus 154 and 129 pg/ml for IL-1Ra, P = 0.03 and 25 versus 9 and 8.8 pg/ml for TNF- $\alpha$ , P = 0.045).

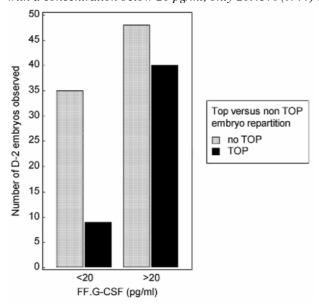
Of 96 embryos observed at the early cleavage stage (25 h), 31 cleaved at 25 h and 65 did not (remaining at either the 2 or 0 pronuclei stage). FF IL-2 and IFN- $\gamma$  were significantly higher in early cleavage embryo compared with the others (10 versus 7.3 pg/ml for IL-2 and 55 versus 24.4 pg/ml for interferon gamma, P = 0.03 and 0.003, respectively). On Day 2, application of the embryo score rated 68 embryos at 16 or higher, 28 from 15 through 12 and 36 below 12. IL-12 was higher in the FF from embryos scored below 12 (17 versus 14 pg/ml in the two other groups, P = 0.05).

Applying the embryo grading, 81 embryos were Grade I, 29 were Grade II, 17 were Grade III and 5 were Grade IV. IL-12 was significantly higher in FF from Grade IV embryos when compared with Grade I, II and III embryos (23.8 versus 14.6, 15.4, 16.23 pg/ml, respectively, P = 0.013).

There were 49 embryos classified as 'Top' embryos and 83 as 'non-Top' embryos. The FF CCL5 levels were significantly higher for those corresponding to high quality 'Top' embryos than for the others (756 versus 271 pg/ml, P = 0.01).

When we compare the group of embryos derived from follicles with an FF G-CSF concentration <20 and  $\ge20$  pg/ml, the ratio of Top to not Top embryos differed significantly (20.45 versus 45%, P = 0007) (Fig. 2).

**Figure 2:** 'Top' versus 'non-Top' repartition in function of FF G-CSF content (pg/ml). In FF with a concentration of G-CSF of 20 pg/ml or higher, 45% (40/88) of the embryos were of 'Top' quality. For the ones with a concentration below 20 pg/ml, only 20.45% (9/44) were 'Top' embryos (P = 0.007).

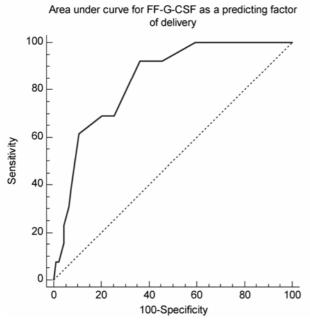


Cytokines, chemokines and implantation/delivery rates

The level of only one of the substances tested—granulocyte-colony stimulating factor (G-CSF)—was correlated with the implantation potential of the corresponding embryos (r = 0.40, P < 0.001). The embryos were classified according to their implantation and delivery rates. Only the embryos with a clear outcome—that is, those which did not implant (n = 89) and those which that definitely implanted and were delivered (n = 13)—were used to construct the AUC-ROC for G-CSF. The area under the ROC curve for ongoing implantation was 0.82 (0.73-0.89) and highly significant (P = 0.0001). The area under the ROC curve for successful delivery was 0.83 (0.75-0.90) and was also highly significant (P = 0.0001) (Fig. 3). We used the AUC-ROC to define an upper and lower limit for G-CSF to determine whether the G-CSF concentration could be used to evaluate the implantation potential for each oocyte and thus for each corresponding embryo, in order to help select which and how many embryos should be transferred. The lower limit was defined according to the AUC-ROC by the highest negative predictive value of implantation (G-CSF < 20 pg/ml). When FF G-CSF was lower than 20 pg/ml, the negative predictive value was 100%. The upper limit was defined by the highest positive predictive value of implantation. According to the AUC-ROC, when the FF G-CSF exceeded 24 pg/ml, the positive predictive value reached 40%.

When we evaluated the 132 transferred embryos according to their FF G-CSF levels, we observed very significant differences in implantation rate between embryos with low and intermediate G-CSF levels (implantation and delivery rates were 9 and 6% compared with 18 and 15.8%, respectively, P = 0.003 and < 0.001) and especially between the embryos with low and high G-CSF levels (9 and 6 versus 44% for both implantation and delivery, P < 0.0001) (Table II).

Figure 3: ROC curve for measurements of G-CSF in individual FF samples from patients with 'No' and 'Certain delivery' groups. The true positive rate (sensitivity) is plotted as a function of the false-positive rate (100-specificity) for different cut-off points of FF-GCSF concentration. Each point on the ROC plot represents a sensitivity/ specificity pair corresponding to a particular decision threshold. AUC-ROC curve is a measure of how well FF G-CSF can distinguish between two main diagnostic groups (certain delivery/no delivery). (Continuous line) The area under curve is 0.84, indicating that a randomly selected individual from the positive group has a test value larger than that for a randomly chosen individual from the negative group 84% of the time. (Dotted lines) The area under the ROC curve is 0.5, representing the null hypothesis.



**Table II.** Implantation and delivery rates as a function of the FF G-CSF concentrations in the individual FF samples corresponding to the fertilised oocytes.

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G-CSF concentrations (Luminex Bio-Rad)	Number of embryos concerned	Mean implantation rate (%)	Mean delivery rate (%)
Low G-CSF ( <20 pg/ml)	45	9	6
Intermediate G-CSF (Between 20 to 24 pg/ml)	62	18*	15.6*
<b>High G-CSF</b> (>24 pg/ml)	25	44**	44**

The lower threshold of implantation was defined by a negative predictive value of 100% from the AUC-ROC curve (<20 pg/ml). The higher threshold of implantation was defined by the highest positive predictive value for implantation (over 24 pg/ml). \*P = 0.0005 and 0.001 between intermediate and low G-CSF for implantation and delivery rates. \*\*P < 0.0001 between high and low G-CSF for both implantation and delivery rates.

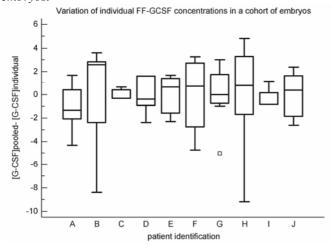
# Variations of [FF]G-CSF within a single ovarian cohort

To determine whether useful information could be obtained by measuring the G-CSF concentration in pooled FF or whether it was essential to measure it for each individual follicle and oocyte, we evaluated the samples from every oocyte producing an embryo among 15 patients, independently of the outcome (embryo transfer, freezing or destruction). This allowed us to compare the variations between samples within a single ovarian cohort, that is, the oocytes produced by a single woman in a single cycle. We evaluated 76 samples with the Luminex kit manufactured by Bio-Rad. The mean value of G-CSF in the pooled FF (evaluated by adding together the separate FF G-CSF measurements for each woman and taking the mean) did not reflect the variations observed

## in individual FF samples.

To assess the variations from the mean in a given ovarian cohort, we determined for each sample the following ratio: mean [FF] G-CSF in the cohort/individual [FF] G-CSF concentration. Fig. 4 shows this variation among 10 patients. Each box shows the variation of individual FF samples from the mean of all the embryos generated by these patients. The results strongly suggest that the embryos generated are not equally correlated with FF G-CSF and differ in their individual implantation potential. It is therefore the individual G-CSF value of the FF sample that reflects its implantation potential. Even in apparently good responders, there are embryos with a low implantation potential and vice versa.

**Figure 4:** The variation in concentration of individual FF in the same cohort of embryos obtained from 10 patients. Each box shows the variation of individual FF samples from the mean in the same cohort of generated embryos.



## Discussion

The measurement of cytokines and chemokines of fluid from individual follicles revealed that the levels of several key immune factors were related to either the morphology of the related embryo or its implantation potential.

We observed that the development of the embryo during the first two days of *in vitro* culture differed significantly in terms of the early cleavage step. Early cleavage at 25 h has been described as a good prognostic factor, when morphology remains good on Day 2 (Lundin *et al.*, 2001; Giorgetti *et al.*, 2007; Sundstrom and Saldeen, 2007; Terriou *et al.*, 2007). Interferon gamma and IL-2 are well-known Th-1 cytokines and are found at significantly higher levels in FF of embryos that cleave early. This suggests that some Th-1 cytokines play a positive role in the process of embryo cleavage. In contrast, the significant elevation of IL-12, known to be cytotoxic, in highly fragmented embryos on Day 2, suggests that it has a deleterious role. This supports the previous finding that elevated IL-12 levels in pooled FFs is associated with a poor pregnancy rate (Gazvani *et al.*, 2002; Bedaiwy *et al.*, 2007). CCL5 was significantly higher in FF from 'Top' embryos. CCL5, which is thought to be produced by the surrounding T cells present in granulosa cells, acts as a chemo-attractant and may play a role. Even though preliminary, these results suggest that the assessment of the levels of some selected cytokines in individual FFs may help embryologists in their assessment of pre-implantation development.

The main finding of this study, however, is that G-CSF in individual FFs appears to be an immune biomarker predictive of successful implantation and delivery. It can be measured easily before embryo transfer and even before fertilization itself, and may therefore be helpful in a strategy of SET. G-CSF is a member of the colony-stimulating-factor family, initially described as a haematopoietic growth factor (Clark and Kamen, 1987). Its main role is to act on the proliferation, differentiation and activation of haematopoietic cells of the neutrophilic lineage (Visani and Manfroi, 1995; Mielcarek *et al.*, 1996). Although produced primarily by the latter, G-CSF is also secreted by other cells, including some in the reproductive tract: human luteinized follicular granulosa cells (Salmassi *et al.*, 2004), endometrial cells (Giacomini *et al.*, 1995), and cells from decidual, placental (Duan, 1990; Miyama *et al.*, 1998) and various foetal (Calhoun *et al.*, 1999) tissues.

Studies using western blot and immunohistochemistry techniques have located the G-CSF protein and its receptor in the ovary—mainly in the granulosa cells of the follicle and in luteal cells (Salmassi *et al.*, 2004). At

ovulation, G-CSF concentrations are much higher in FF than in serum (Salmassi *et al.*, 2005). This observation is quite common for the cytokines, chemokines, hormones and growth factors involved in the complex process of ovulation.

Previous studies of G-CSF in serum provided some evidence of its involvement in implantation: some authors reported that its serum levels rise at implantation in the case of successful natural cycles (Yanagi *et al.*, 2002) and after successful IVF/ ICSI attempts (Salmassi *et al.*, 2005). The source of this secretion seems to be the decidual and endometrial cells in the uterus, which indicates that it occurs about 10 days after oocyte retrieval (Duan, 1990; Giacomini *et al.*, 1995).

Low levels of FF G-CSF were observed in ageing patients and in high responders to stimulation, both characteristics known to be associated with decreased pregnancy rates and to affect oocyte quality. Thus indirectly, we may postulate that the FF G-CSF is linked to oocyte quality and could represent a new tool for the evaluation of individual oocyte competence. The mean FF G-CSF concentration in women older than 36 was below the threshold of 20 pg/ml, the category associated with the highest negative predictive value for implantation. In such patients, non-invasive tools able to identify the oocyte with the best potential would be of capital importance.

Significantly more 'Top' embryos were obtained when the FF G-CSF was 20 pg/ml or higher, even though FF-G CSF levels were not different between 'Top' and 'not Top' embryos. In terms of a strategy to choose which embryo to transfer, we may postulate that FF G-CSF and morphological development of the pre-implantation embryo are complementary tools that would not lead to discrepant evaluations. Assessment of other cytokines, such as CCL5 (for a positive selection) or IL-12 (for a negative selective) may easily complement the analysis if necessary. We also should not forget that new non-invasive tools focusing on the embryo itself are developing and could be a useful complement (Sakkas and Gardner, 2005; Seli *et al.*, 2007; Scott *et al.*, 2008).

The assessment of G-CSF in individual FFs 2 days before embryo transfer may be very useful in cohorts in which all the embryos appear equal according to morphological criteria. This information would also be valuable in selecting which embryo(s) to freeze.

How [FF]G-CSF influences future embryo implantation is still unknown. However, there are several possibilities.

First, the FF itself may be involved in the complex dialogue that leads to implantation and could transmit messages directly to the fallopian tube and to the uterus in fertile patients (Leese *et al.*, 2008). If we consider pregnancy as a semi-allograft, the question of maternal immune tolerance is essential. It has been reported that pretreatment of mice with G-CSF before an allograft promotes T cell tolerance towards these grafts through the induction of a switch towards a Th-2 dominated response and tolerogenic dendritic cell differentiation towards DC2 (Pan *et al.*, 1995). Furthermore, G-CSF promotes the generation of IL-10-producing T-regulatory (T-reg) cells and promotes transplantation tolerance (Morris *et al.*, 2004), a feature obviously relevant to the immunoregulatory events occurring in the pre-, peri- and early post-implantation uterus. Thus, we may speculate that the [FF]G-CSF level reflects the early oocyte-uterus crosstalk that leads to the preparation of a receptive uterus, even before the embryo is generated (Herrler *et al.*, 2003). One criterion for each oocyte's quality might therefore be its intrinsic capacity to prepare the uterus and to engage it in an immune tolerance-promoting pathway.

A second hypothesis is that [FF]G-CSF influences the mRNA content of the oocyte itself. In particular, it may induce adhesion molecules (including L selectin) at the oocyte cell surface to promote the adhesion of the future embryo to the endometrial cells. Alternatively, G-CSF could induce the cumulus granulosa cells to produce the cytokines or growth factors which are necessary for the development and implantation of the resulting embryo. It has been reported that both macrophages and CD4+T cells are normally detectable in the cumulus oophorus. More importantly, these T cells produce higher levels of IL-4 and LIF than the T cells of peripheral blood or ovarian cells isolated from the same women (Piccinni *et al.*, 2001). Thus, [FF]G-CSF, which can induce a Th2 switch, (Rutella *et al.*, 2005) could be responsible, at least in part, for IL-4 production by the cumulus oophorus T cells, which in turn induces the production of LIF by T cells (Piccinni *et al.*, 1998). Amongst its many properties related to implantation, LIF has positive effects on the growth and differentiation of preimplantation human blastocysts and significantly increases the quality and number of human blastocysts *in vitro* (Dunglison *et al.*, 1996).

Another hypothesis is that [FF]G-CSF provides the embryo with crucial information on how to repair itself. Widespread use of pre-implantation diagnosis has shown not only the high incidence of mosaicism and aneuploidy in embryos in culture (Baart *et al.*, 2006), but also their self-repair (Munne *et al.*, 2005). Self-repair has been described in haematopoietic stem cells and appears to be affected by ageing (Rossi *et al.*, 2007), as also observed in embryos. G-CSF has been described in various models (heart, liver) as an agent promoting

endogenous repair (Yannaki *et al.*, 2005) either through the mobilization of haematopoietic multipotent progenitor cells (Imamura *et al.*, 2005) or the enhancement of endogenous stem cells *per se* (Piscaglia *et al.*, 2007).

All these hypotheses need further detailed investigations with animal models before their eventual validation in humans.

To conclude, although a prospective randomized study is essential to validate our hypothesis, this preliminary report strongly suggests that assessing the concentration of G-CSF in individual FF may be a valuable, non-invasive method to select those oocytes (and hence the embryos subsequently generated from them) with the highest implantation potential. It may also be a valuable biomarker to evaluate implantation potential among patients with a low ovarian reserve, which is still the major limiting step in our daily practice. In a large cohort, it would help the embryologist to choose which embryo to transfer or to freeze. This should make it possible to increase implantation rates significantly and facilitate the universal adoption of a 'SET' policy, without decreasing the subsequent pregnancy rates. It would therefore make it possible to reduce multiple pregnancy rates in IVF/ICSI-ET cycles and their related morbidity and mortality.

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## **Author Roles**

N.L. designed the clinical setting, organized the individual FF collection and analysed the results. MP Piccinni designed the multiplex cytokine analysis, chose the cytokines to be tested, performed the Luminex analysis, with the help of L.L. R.L. and J.S. were the head of the clinical and biological Reproductive Department. Y.V. was the head of Obstetrics and Gynaecology department at Poissy Hospital. S.D. organized the shipment of samples and ensured their traceability. G.C., F.F. and J.M.F helped to decide on the cytokines and chemokines to be assessed and helped to interpret the results. Prof. S.R. was the director of the DENOTHE Centre of Excellence for Research, Transfer and Higher Education at the University of Florence and of the Department of Internal Medicine (Florence, Italy). Prof. E.M. was the director of the Immunoallergology unit (Florence, Italy). We applied for a provisional patent July, 2006 and a PCT patent of application July, 2007 (WO-2008/009705, accepted 24.01.08).

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