

USER MANUAL

Ampliquality CHLAM-T

Code 03-10A Code 03-10R

Kit for the detection of Chlamydia trachomatis





1.	PRODUCT INFORMATION	3
2.	KIT CONTENTS	4
3.	STORAGE AND STABILITY OF THE REAGENTS	5
4.	PRECAUTIONS FOR USE	6
5.	SAFETY RULES	7
5.1.	General safety rules	7
5.2.	Safety rules about the kit	8
6.	MATERIALS REQUIRED, BUT NOT PROVIDED	9
6.1.	Reagents	9
6.2.	Instruments	9
6.3.	Materials	9
7.	PREPARATION OF THE REAGENTS	10
8.	INTRODUCTION	11
9.	TEST PRINCIPLE	12
10.	PRODUCT DESCRIPTION	13
11.	COLLECTION, MANIPULATION AND PRE-TREATMENT OF THE SAMPLES	.14
12.	PROTOCOL	15
12.1.	DNA extraction	15
12.2	DNA amplification 2.1. TST DNA amplification 2.2. Direct amplification (1 st amplification) of C. trachomatis DNA:	15 15 15
12.3.	I I	
	3.1. Agarose gel electrophoresis 3.2. Sample Loading	16 17
12.4. ampli	Interpretation of results: TST DNA amplification and C. trachomatis first	18



12.5.	2 nd C. trachomatis DNA amplification (nested PCR)	19
12.6.	Interpretation of results: 2 nd amplification of Chlamydia trachomatis DNA	۹.20
13.	TROUBLESHOOTING	21
14.	DEVICE LIMITS	23
15.	DEVICE PERFORMANCES	23
15.1.	Specificity	23
15.2.	Sensitivity	23
16.	REFERENCES	24
16.1.	Useful websites	24



1. PRODUCT INFORMATION

CHLAM-T:

Kit for the detection of *Chlamydia trachomatis* by DNA amplification in the cryptic plasmid.

The kit contains the reagents for DNA amplification and visualization by agarose gel electrophoresis, the internal control of sample amplificability, and positive control.

Code	Product	Pkg
03-10A-25	CHLAM-T	25 tests
03-10A-50	CHLAM-T	50 tests

CHLAM-T - amplification reagents

Kit for the detection of *Chlamydia trachomatis* by DNA amplification in the cryptic plasmid.

The kit contains the reagents for DNA amplification, the internal control of sample amplifability, and the positive control.

Code	Product	Pkg
03-10R-25	CHLAM-T- amplification reagents	25 tests
03-10R-50	CHLAM-T- amplification reagents	50 tests



2. KIT CONTENTS

Attention: the different contents correspond to different kits.

(legend: **X** = component included in the kit; **0** = component not included in the kit)

BOX P

STORE AT - 20°C

code 03-10A	code 03-10R	DESCRIPTION	LABEL	COLOUR OF TUBE (T) OR LID	25 tests	50 tests	8 tests
X	X	Single dose premixes for CHLAM-T direct amplification		Colourless (T)	25	50	8
х	X	Single dose premixes for CHLAM-T nested amplification		Green (T)	25	50	8
X	x	Single dose premixes for TST (thiosulfate sulfurtransferase –rhodanese) amplification		Blue (T)	25	50	8
Х	х	Thermostable Taq DNA polymerase	AB TAQ 5 U/μL	Red	1 x 50μL	1 x 100μL	1 x 20μL

SMALL BAG

STORE AT - 20°C

code 03-10A	code 03-10R	DESCRIPTION	LABEL	COLOUR OF TUBE (T) OR LID	25 tests	50 tests	8 tests
x	x	Plasmid DNA containing part of <i>Chlamidia trachomatis</i> genome	CHLAM-T positive control	Blue	1 x 30 μL	1 x 50 μL	1 x 10 μL

BOX F

STORE AT +2°/ +8°C

code 03-10A	code 03-10R	DESCRIPTION	LABEL	COLOUR OF TUBE (T) OR LID	25 tests	50 tests	8 tests
x	0	Loading Buffer for Electrophoresis	6X Blue	Blue	1 X250μL	1 X450μL	1 X100μL
Х	0	Ethidium Bromide solution (2,5 mg/mL)	Ethidium Bromide TOXIC R 23 68 S 36/37 45	Red	1 X170μL	1 X250μL	1 X100μL
Х	0	DNA Molecular Weight Marker	MW Marker	Yellow	1 X200μL	1 X400μL	1 X70μL



BOX A

STORE AT +15°/ +25°C

code 03-10A	code 03-10R	DESCRIPTION	LABEL	COLOUR OF TUBE (T) OR LID	25 tests	50 tests	8 tests
X	0	Agarose for Molecular Biology	AGAROSE		1X20g	1X40g	1X6g
X	0	Electrophoresis buffer TRIS-Acetate-EDTA, pH 8,00	50X TAE		1X60mL	1X125mL	1X30mL

3. STORAGE AND STABILITY OF THE REAGENTS

Each component of the kit must be stored according to the directions indicated on the label of the single boxes. In particular:

Box P	store at -20°C
Small bag	store at -20°C
Box F	store at a +2°- +8°C
Box A	store at +15°- +25°C

When stored at the recommended temperature, all test reagents are stable until their expiration date.



4. PRECAUTIONS FOR USE

- The kit must be handled by qualified investigators, who are educated and trained in molecular biology techniques applied to diagnostics;
- Before starting the kit procedure, read carefully and completely the instruction manual;
- Keep the product away from heat sources;
- Do not use any part of the kit past the expiration date;
- In case of any doubts or questions about the storage conditions or box integrity, contact AB ANALITICA's technical support at: laboratorio@abanalitica.it;

In the amplification of nucleic acids, the investigator has to take the following special precautions:

- Use filter-tips;
- Store the biological samples, the purified DNA, the reference DNA included in the kit, and all the amplification products in different places from where amplification reagents are stored;
- Organize the work space into different pre- and post-PCR units; do not share instruments and consumables (pipettes, tips, tubes, etc.) between them;
- Change gloves frequently;
- Wash the bench surfaces with 5% sodium hypochloride;
- The PCR premixes are thawed at room temperature before use; Add the Taq DNA polymerase and purified DNA very quickly at room temperature or in an ice-bath;



5. SAFETY RULES

5.1. General safety rules

- Wear disposable gloves to handle the reagents and the clinical samples and wash your hands at the end of work;
- Do not pipette with the mouth;
- Since, no known diagnostic method can assure the absence of infective agents. It is a good rule to consider every clinical sample as potentially infectious and handle it as such;
- All the devices that directly touch the clinical samples must be considered as contaminated and disposed as such. In case of accidental spilling of the samples, clean up with 10% Sodium Hypochloride. The materials used to clean up must be disposed of in special containers for contaminated products;
- Clinical samples, materials, and contaminated products should be disposed of after decontamination by:

immersion in a solution of 5% Sodium Hypochloride (1 volume of Sodium Hypochloride solution for every 10 volumes of contaminated fluid) for 30 min.

OR

autoclave at 121°C for at least 2 hours (NOTE: do not autoclave solutions containing Sodium Hypochloride!!)



5.2. Safety rules about the kit

The risks for the use of this kit are related to the single components:

Dangerous components:

ETHIDIUM BROMIDE (included in the 03-10A kit)

3,8-diamino-1-ethyl-6-phenylphenantridiumbromide <2%

Description of risk: T (Toxic)

RISK SENTENCES AND S SENTENCES

ETHIDIUM BROMIDE

R 23 and R 68 Toxic for inhalation.

Risk of irreversible effects.

S 36/37 45 Wear laboratory coat and disposable gloves.

In case of accident or discomfort, seek for medical

assistance and show the package label.

R and S sentences refer to the concentrated product, as provided in the kit. In particular for Ethidium Bromide, until it is diluted in the agarose gel.

When handling concentrated Ethidium Bromide, use a chemical dispensing fume cabinet. Always wear disposable gloves and laboratory coat when handling diluted Ethidium Bromide solutions as well.

The product must **not** be disposed of in the common waste. It must not go in the drain system. For disposal, follow local laws.

In case of accidental spilling of Ethidium Bromide, clean with Sodium hypochloride and water.

Material Safety data sheets of products are available upon request.



6. MATERIALS REQUIRED, BUT NOT PROVIDED

6.1. Reagents

- Reagents for DNA extraction;
- Sterile DNase and RNase free water;
- Distilled water:
- Reagents for DNA visualization by agarose gel electrophoresis (necessary for 03-10R kit).

6.2. Instruments

- Laminar flow cabinet (use is recommended while adding TAQ polymerase to the amplification premix to avoid contamination; it would be recommended to use another laminar flow cabinet to add the extracted DNA);
- Micropipettes (range: 0,2-2 μL; 0,5-10 μL; 2-20 μL);
- Thermalcycler;
- Microcentrifuge (max 12.000-14.000 rpm);
- Balance;
- Magnetic heating stirrer or microwave;
- Chemical cabinet (use is recommended in handling Ethidium Bromide);
- Horizontal electrophoresis chamber for agarose minigel;
- Power supply (50-150 V);
- UV Transilluminator;
- Photo camera or image analyzer.

6.3. Materials

- Disposable gloves;
- Disposable sterile filter-tips (range: 0,2-2 μL; 0,5-10 μL; 2-20 μL;);
- Graduated cylinders (1 L) for TAE dilution;
- Pyrex bottle or Becker for agarose gel preparation;
- Parafilm.



7. PREPARATION OF THE REAGENTS

Preparation of 1 L of 1X TAE buffer:

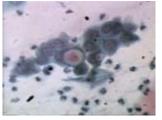
Mix 20 mL of 50X TAE (included in the 03-10A kit) with 980 mL of distilled water.



8. INTRODUCTION

Chlamydia trachomatis is the ethyologic agent of several common sexually transmitted bacterial pathologies.

Chlamydia genera encompasses low-dimension microorganisms, that



reproduce exclusively in the cytoplasm of the host cell, in the vacuole. It is a bacterium with obliged endo-cellular parasitism (Fig. 1), due to its incapability to produce ATP and other necessary biosynthetic cofactors.

Its morphogenetic cycle consists of two alternate cellular cycles: the **elementary body** and the **reticular body**.

Cytoplasmic inclusion of Chlamydia trachomatis

The elementary body is the infective form and, for its survival capacity outside the eukaryotic cell, can be compared to a spore cell. Once it penetrates the eukaryotic cell, the elementary body develops after 10 hours from the infection. Then, the reticular body starts to reproduce up to cell death.

Chlamydia trachomatis encloses 15 distinct serotypes: A,B,C serotypes are associated to trachoma (conjunctive infection); L1, L2, and L3 serotypes are associated to lymphogranuloma venereum; D to K serotypes are ubiquitous, generally associated to sexually-transmitted diseases; they infect the epithelial cells and cause infection in the pharynx, respiratory apparatus, urethra, and uterine tubes (also called Fallopian tubes).

Each serotype has a plasmid (about ten copies per cell) that is believed to contain the essential genes for microorganism survival in the infected cell.

Generally, *C. trachomatis* infection has a clinically latent form but can be the cause of ectopic pregnancy, female sterility, and male infertility. Newborns exposed to C. trachomatis infection can develop a conjunctivitis or an acute pneumonia that can lead to a chronic form.

Most of the *C. trachomatis* infections are asymptomatic and it is generally difficult to distinguish an evident C. trachomatis infection from other sexuallytransmitted diseases; therefore it is important to have a quick, rapid, and reliable method for C. trachomatis detection.

Recent studies suggest that there is a correlation between the exposure to different C. trachomatis serotypes and an increased risk of cervix cancer (Antilla et al. JAMA 2001).



Even if HPV infection from oncogenic strains is the first cause of cervix cancer, the researchers are now focusing on *C. trachomatis*.

It is not clear if this oncogenic risk is associated to specific microorganism's serotypes (Koskela *et al.*, 2000).

Considering the high incidence (4%) of *C. trachomatis* infection in male donors, guidelines for the prevention of this pathogen's transmission by artificial insemination have been recently proposed.

The American Society for Reproductive Medicine and the European Society for Human Reproduction and Embryology affirm that tests for *C. trachomatis* identification should always be performed to screen urethral, urine, and seminal samples (American Society for Reproductive Medicine 1998, Baratt et al.,1998).

9. TEST PRINCIPLE

PCR (**P**olymerase **C**hain **R**eaction) was the first method of DNA amplification described in literature. (Saiki RK et al., 1985). It can be defined as an *in vitro* amplification reaction of a specific part of DNA (target sequence) by a thermo-stable DNA polymerase.

In the reaction, three segments of nucleic acid are involved: the double strand DNA template to be amplified (target DNA) and two single-strand oligonucleotides "primers" that anneal in a specific way to the template DNA. The DNA polymerase begins the synthesis process at the region marked by the primers and synthesizes new double stranded DNA molecules, identical to the original double stranded target DNA region, by facilitating the binding and joining of the complementary nucleotides that are free in solution (dNTPs). After several cycles, we can get millions of DNA molecules which correspond to the target sequence.

The sensitivity of this test makes it particularly suitable to the application in laboratory diagnostic.

Besides, the amplification reaction can be executed from a wide range of biological samples, and since PCR is able to amplify very small DNA segments, the starting DNA can also be partially degraded.



10. PRODUCT DESCRIPTION

The method proposed in this kit consists of the amplification of the *C. trachomatis* cryptic plasmid DNA.

The plasmid is highly conserved in all sierotypes of *C. trachomatis*, but absent in other species.

The method can evaluate the suitability of extracted DNA, by amplification of TST (thiosulfate sulfurtransferase rhodanese) gene in the region 22q13.1 (amplification control). A negative result in the TST gene amplification indicates either the presence of inhibitors of the amplification reaction in the extracted DNA or that the DNA is highly degraded. This method helps the operator to recognize possible false negative results for *C. trachomatis*.

The kit also provides a positive control for the amplification. When the amplification of the positive control is successful, this guarantees that the reaction worked correctly. The positive control is not dangerous for the operator, because it is plasmid DNA containing only a part of *C. trachomatis* cryptic plasmid.

The kit is in premix format: all the reagents for the amplification are pre-mixed and aliquoted in single dose test-tubes, which Taq polymerase and the extracted DNA will be added to. This premix format reduces the manipulation steps in pre-amplification, saves considerable time for the operator, avoids repeated freezing/thawing of reagents (that could alter the product performances) and, above all, this format reduces significantly the risk of contamination and the risk of obtaining false positive results. For this reason, it is always recommended to use all the proper amplification controls.



11. COLLECTION, MANIPULATION AND PRE-TREATMENT OF THE SAMPLES

The samples used in the laboratories for the determination of *C. trachomatis* infection, are cytological, like cell cultures, urethral swabs and cervix-vaginal swabs, sperm, urine, etc..

In the case of cervix and vaginal swabs, sample collection is performed by using a little swab or a little sterile brush (*cyto-brush*). The collected cells are diluted in a suitable transport media (1X PBS, physiological solution or other medium).

The sample can be stored at $+2 / +8^{\circ}$ C for a max of 48 hrs.; after that, it is better to proceed with nucleic acid extraction.

If it is not possible to extract the sample within a short period of time, the specimen must be stored at -20°C.

When sample cellularity is too low, it could be necessary, before starting DNA extraction, to concentrate the sample by centrifuging repeatedly. For sample resuspension use sterile PBS, which can also be useful to remove mucus and red blood cells, if present.

Seminal liquid is collected in a sterile container and stored at +4°C for some hours (maximum above night). Alternatively, it is frozen by adding a specific medium, which compatible with the DNA extraction method used in the molecular analysis.

C. trachomatis detection from urine samples is usually performed from the first urination. It is necessary to discard the first part of urine flux and to collect only the second one in a sterile container.



12. PROTOCOL

12.1. DNA extraction

Any extraction method can be used, provided that it extracts pure and integral *C. trachomatis* DNA.

AB ANALITICA tested successfully methods based on spin columns (e.g. QIAmp DNA mini kit - QIAGEN) and methods based on matrixs (e.g. INsta-Gene Matrix – BIO-RAD).

It is necessary to remember that the volumes of the extracted DNA, indicated in paragraph 12.2 for TST and *C. trachomatis* amplification, are referred to the extracts obtained with AB ANALTICA methods. When using an alternative extraction method, it will be necessary to adjust the volume of the amplification mix by adding water if the volume of the DNA solution for amplification is less than 20 μL .

12.2. DNA amplification

12.2.1. TST DNA amplification

For each sample add to each premix tube (blue tube):

0,5 μL AB Taq
10 μL extracted DNA

12.2.2. Direct amplification (1st amplification) of C. trachomatis DNA:

For each sample add to each premixed colourless tube:

0,5 μL AB Taq
10 μL extracted DNA



It is important to include in each experiment a negative control (add 10 μ L of distilled H₂O instead of DNA to the mix) and a positive control (10 μ L of the positive control included in the kit).

Centrifuge briefly, put the tubes (TST and CHLAM-T) into the thermalcycler as programmed below:

1 cycle	95°C	5 min
	95°C	45 sec
30 cycles	56°C	45 sec
	72°C	45 sec
1 cycle	72°C	5 min

Amplification products length:

CHLAM-T: 594 bp. TST: 202 bp.

12.3. Visualization of the amplification products

12.3.1. Agarose gel electrophoresis

Prepare a 3% agarose gel:

Weight 1.5 g of agarose and pour it into 50 mL of 1x TAE.

Leave the solution on a magnetic stirring heater or in a microwave, until the solution becomes clear.

Allow the gel to cool for a few minutes and then add 10 μ L of Ethidium Bromide solution.

NOTICE: Ethidium Bromide is a strong mutagenic agent; always wear gloves and preferably work under a chemical safety cabinet during the handling of this reagent or gels containing it.

Pour the gel into the appropriate gel casting tray, with the comb placed in and allow the gel to cool at room temperature or in a fridge until the gel becomes solid.



When the gel is solidified, remove the comb carefully (pay attention not to damage the gel wells) and transfer the tray into the electrophoresis chamber. Pour the appropriate amount of 1x TAE buffer, so that it covers the gel completely (about 1-2 mm over the gel surface).

12.3.2. Sample Loading

Mix in a tube or directly on a parafilm layer:

2 μL of 6X Blue*

10 μL of amplification product or DNA Molecular Weight Marker

(MW*marker)

Load the mixture in the gel wells; switch on the power supply and set the voltage between 80-100 V.

Run the gel for about 40 min, then place the gel on an UV transilluminator and analyze the results by comparing the size of the amplification products with the reference Molecular Weight Marker.

* = 6X Blue and PM Marker are included in 03-10A kit; for the use of another loading buffer or of another Molecular Weight Marker, follow the indications specified by the supplier.

DNA Molecular Weight Marker (MW Marker, included in 03-10A kit)

Fragment sizes: 501-489, 404, 331, 242, 190, 147, 111, 110, 67, 34x2, 26 bp.

The 501-489 bp bands usually are not clearly distinguishable and appear as an unique band; the 26 and 34 bp bands are sometimes too small to be seen in a 3% agarose gel (because of their low molecular weight).

NOTICE: UV rays are dangerous for skin and, above all, eyes: always wear gloves and safety glass or use the protection screen of the UV transilluminator.



12.4. Interpretation of results: TST DNA amplification and C. trachomatis first amplification

The controls should show the following results:

CONTROL	RESULT	INTERPRETATION
Positive control		The PCR amplification was performed correctly
Negative control	NEGATIVE	Absence of contamination

Then, for the interpretation of the band pattern, follow the table below:

RESULT	INTERPRETATION		
Absence of the TST gene band	Sample not suitable for amplification (repeat the DNA extraction)		
Presence of the TST gene band Absence of <i>Chlamydia trachomatis</i> band	Amplifiable sample Proceed with the second amplification to evaluate the real negativity of the sample		
Presence of the TST gene band Presence of <i>Chlamydia trachomatis</i> band	Amplifiable and positive sample for <i>Chlamydia trachomatis</i>		



12.5. 2nd C. trachomatis DNA amplification (nested PCR)

Add to each premixed green tube, containing 48.5 µL of amplification mix:

0,5 µL AB Taq

1 μL First amplification product

It is important to use in this step the negative and the positive controls included in the *Chlamydia trachomatis* direct amplification. When it seems useful, it is possible to include also a negative control for the nested amplification, by adding 1 μ L of sterile water to the nested mix, instead of the direct amplification product.

Centrifuge briefly, then put the tubes into the thermalcycler as programmed below:

1 cycle	95°C	5 min
	94°C	45 sec
35 cycles	55°C	45 sec
	72°C	45 sec
1 cycle	72°C	10 min

Amplification fragment length: 201 bp.

At the end of the amplification program, visualize the nested amplification products by a 3% agarose gel electrophoresis.

Weight 1.5 g of agarose and pour it into 50 mL of 1X TAE.

Follow the instructions indicated at paragraph 12.3.



12.6. Interpretation of results: 2nd amplification of Chlamydia trachomatis DNA

The controls should show the following results:

CONTROL	RESULT	INTERPRETATION		
Positive control	POSITIVE	The first and the second amplification was performed correctly		
Negative Control	NEGATIVE	Absence of contamination in the first amplification		
Negative control of Nested amplification	NEGATIVE	Absence of contamination in the second amplification		

Then, for the interpretation of the band pattern, follow the table below:

RESULT	INTERPRETATION
Absence of Chlamydia trachomatis band in	
the first and second amplification	Chlamydia trachomatis
Presence of Chlamydia trachomatis band in	Positive sample for
the second amplification	Chlamydia trachomatis

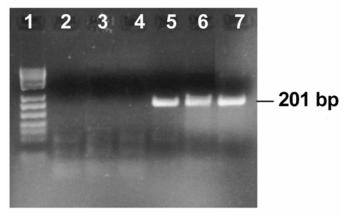


Fig. 2: 3% agarose gel electrophoresis of the amplification products.

- 1. DNA Molecular Weight Marker.
- 2 4. Direct PCR products.
- 5-7. Nested PCR products.

All samples are positive for *C. trachomatis*.



13. TROUBLESHOOTING

No amplification products or positive control DNA band

- TAQ polymerase was not correctly added to the premix
 - Use pipettes and tips with suitable volumes (pipette range 0,2 2 μL);
 - Check visually that TAQ polymerase diffuse in the premix: this is easy because the enzyme is in dissolved glycerol that has a higher density;
 - Alternatively, check visually the drop of TAQ polymerase put on the tube wall, then centrifuge briefly.
- The thermalcycler was not programmed correctly.
 - Check the conformity of the thermalcycler program and the temperature profile in the instruction manual.
- The kit does not work properly
 - Store the premix, enzymes and reference DNA at -20°C;
 - Avoid repeated freezing/thawing of the premix and the regents.

No amplification bands for either TST or C. trachomatis in the tested sample, but a good band for the Positive controls

- Problems during the extraction step
 - Be sure that the extraction kit is adequate and that you follow correctly all the instructions;
 - Consult the troubleshooting section of the extraction kit;
 - Repeat the DNA extraction starting from a new sample.

The amplification was inhibited

- Dilute the starting sample with distilled water and TE;
- Repeat the DNA extraction from a smaller amount of clinical sample;
- Use an adequate extraction system.



Aspecific products or extrabands present after the visualization of the amplified product in agarose gel

The thermalcycler makes temperature changes too slowly

- Execute a thermalcycler revision.

The preparation of the amplification reaction took too long time at room temperature.

Accelerate the work time at room temperature;

- Work on ice.
- The extracted product was not purified
 - Use an extraction system, which purifies the sample well.

The starting sample contained degraded DNA

- Repeat the extraction step using another starting clinical sample;
- Be sure that the sample had been collected and stored in appropriate way.

For any further problems contact AB ANALITICA's technical support at: laboratorio@abanalitica.it, fax (+39) 049-8709510, or tel. (+39) 049-761698.



14. DEVICE LIMITS

The kit can have reduced performances if:

- the clinical sample is not suitable for this analysis (incorrect transport media or incorrect sample storage);
- the DNA is not amplifiable because of the presence of inhibitors of the amplification reaction or due to an inadequate extraction system;
- the kit was not stored at the suggested temperature.

15. DEVICE PERFORMANCES

15.1. Specificity

Primer sequence alignment in the most important databanks shows the absence of unspecific alignment and guarantees the amplification of *C. trachomatis*. Cross reaction with genomic DNA or other pathogenic microorganism nucleic acids have not been revealed.

15.2. Sensitivity

It has been shown that a unique amplification step, using nested primers, has a detection limit of about 1 IFU (inclusion forming unit) after visualization by agarose gel electrophoresis.

Each IFU contains about 10 cryptic plasmids.

The addition of another amplification step increases the sensitivity by at least a factor of 10.



16. REFERENCES

American Society for Reproductive Medicine. Fertil Steril 4(3), 1S-13S, 1998.

Antilla T, Saikku P, Koskela P et al. JAMA 285(1), 47-51, 2001.

Baratt C, Englert Y, Gottlieb C and Jouannet P. Hum Reprod 13(2), VII-IX, 1998.

Koskela P, Antilla T, Bjorge T et al. Int J Cancer 85, 35-39, 2000.

Saiki RK, Scharf S, Faloona F et al. Science 230, 1350-1354, 1985.

16.1. Useful websites

http://www.drorlandella.it/chlamydia.htm





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